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The invention describes a nucleic acid sequence (I) encoding an Ostertagia ostertagi protein, or part of the nucleic acid sequence encodes an immunogenic fragment of the protein, where the nucleic acid sequence or its part has at least 85% homology with the nucleic acid sequence of the O. ostertagi gene having any of the 7 fully defined sequences of 306-1761 base pairs, as given in the specification. The composition and methods are useful for diagnosing, preventing or tro

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MSAAVVVAV--LLALFSYAEAGFCCPNSLSQSDSARQIFLDFHNDVRRNIALGNGLINWT
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                                              Conservative
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Pred.
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Key Peptide Ancylostoma Z-Z 17 ₩ O Ancylostoma Hawdon Protein 10 10 (UYYA) cylostoma -JAN-1997 W04321; W04321 PSDB; AAT38466. -APR-1995; -APR-1996; 9632479-A1 -OCT-1996. , K UNIV YALE standard; protein; 424 AA. secreted protein; caninum **Becreted** Hotez (first entry) 96WO-US004821. Location/Qualifiers /label= label= Sig_peptide 00419414 protein ASP-1 Jones Mat_protein BF; ASP-1; (pro-form). hookworm; vaccine

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RESULT 4
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virucide; tr
hookworm; in
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Similarity 30.9%;
6; Conservative
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N-PSDB; ADY37245.
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                                                                                                                                                                                                                                                                                                                                                           quence 424 AA;
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Similarity 30.9%;
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                                                        YKVCGTK--LAVSCIYNGVGYITNQPMWETGQACKTGADCSTYKNSGCEDGLCTKGPDVP
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n A, Jones B,
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Pred. No. 4.2e-19;
35; Mismatches 97;
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Hookworm

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23-NOV-2001;
26-APR-2002;
17-OCT-2002;
26-SEP-2003;
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hookworm;
tetanus;
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Loukas
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                                                                                                                                                                                                                                                                                                                           Mendez
                                                                                                                                                                                                                                                  Disclosure;
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WANG Y.
HAWDON J.
LOUKAS A.
WILLIAMSON
JONES B.
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ASHCOM J.
BDAMCHIAN
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MENDEZ S.
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immune response; HIV infection;
diphtheria; pertussis; polio.
                                                                                                                                                                                                                                                                                                                                                                                COUD G
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                                                                       21.9%;
Similarity 30.3%;
74; Conservative
              GLINWTVNADAVILGPAQNMYKVDWDCNLEEVAAQQIAPCNDPLPINTSLAQNIARWLY-
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A, Jones
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FSYAEAGFCCPNSLSQSDSARQIFLDFHNDVRRNIALG
                                                                                                                                                                                                                                                   227pp; English.
                                                                       Score 274; DB 9 Pred. No. 8.6e-17; Mismatches
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G, Bottazzi
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antigen o
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ne invention relates to a composition (I) comprising a copy combinant or synthetic antigen or their fragments derived for a carrier. (I) is useful for vaccinating or eliciting an

from hookworm

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Mendez S;
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                                              Composition useful for vaccinating or eliciting immune response hookworm in mammal, comprises copy of recombinant or synthetic atheir fragments derived from hookworm, and carrier.
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3-NOV-2001;
6-APR-2002;
7-OCT-2002;
6-SEP-2003;
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WILLIAMSON
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BOTTAZZI M
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VCGTK--MATSCIYNGIGYITNAPMWETGQACKTGADCSTYKNSGCEDSLCTKGADVPET
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Williamson
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2001US-0332007P.
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2003US-0505848P.
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1 A, Jones B,
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(HOTE/)
(ASHC/)
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(ZHAN/)
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                                                                                                                                                                                                      antihelmintic; anti-HIV; antitubercular; tuberculostatic; antimalarial virucide; tranquilizer; antibacterial; vaccine; recombinant antigen; hookworm; immune response; HIV infection; tuberculosis; malaria; measl-tetanus; diphtheria; pertussis; polio; gene; ds.
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23-NOV-2001;
26-APR-2002;
17-OCT-2002;
26-SEP-2003;
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HAWDON J.
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WILLIAMSON
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GOUD G.
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ASHCOM J.
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Loukas A,
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                                     nom antigen protein; VAP-1; VAP-2; anaemia; malnutrition; lmonary disorder; nutritional disorder; hyperinfection; nematocide; ichinosis; onchocerciasis; river blindness; lymphatic filariasis; rasitic nematode; antihelmintic; transgenic; chromodosome IV.
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Similarity 31.0%; Pred. No. 1.3e-17;
6; Conservative 34; Mismatches 96
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MENDEZ S.
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                                 PNEVIWQEGKACV-
                                                                   -GEEKGNE-DQKNRFYARRNYFGWANMAKGKTYRVGCSYIMCGDGESALFTCLYNEKAQC
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Query Match Best Local S Matches 62

17.3%; Similarity 29.1%; 2; Conservative

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Score 216.5; DB 5; Pred. No. 6.8e-13; mismatches 77;

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  I; 2002-590738/
PSDB; ABK90244.
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WPI; 2002-N-PSDB; AF

2-590738/63 ABK92046.

Westlund

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Burnam

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Sluder

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18-JAN-2001;

2001US-0263081P

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CAMBRIA BIOSCIENCES

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WO200257440-A2

of a nematode having a tragents for treating, preventing or plants or individuals.

transgene that reguluseful for screening reducing nematode

hat regulates the expre screening anti-nematode

the expression

infestation

Claim 83;

Fig

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105pp;

English

The invention relates to a transgenic nematode, the cells of which contain a transgene comprising a regulatory element of a gene that encodes a nematode secretory product or its homologue operably linked to a DNA sequence encoding a detectable marker. Also included are (1) identifying a compound that inhibits a nematode secretion pathway, for use as an anti-nematode agent for use in preventing or reducing mematode infestation of a plant (the agent may be used to treat the soil or seed); (2) a vector comprising; (a) a vap-1 or vap-2 polymucleotide encoding VAP-1 or VAP-2; (iii) has a sequence comprising the Caenorhabditis elegans vap-1 or vap-2 promoter operably linked to a polymucleotide encoding a detectable marker; or (b) a regulatory element comprising to the C. elegans vap family of genes operably linked to a polymucleotide encoding a detectable marker; (a) expressing the Caenorhabditis of c. elegans amphid sheath cell; and (4) expressing a first polymucleotide in a C. elegans amphid sheath cell; and (4) expressing a polypeptide in a C. elegans amphid sheath cell; and (4) expressing a polypeptide in a C. elegans amphid sheath cell; and (4) expressing a polypeptide in a C. elegans amphid sheath cell; and (4) expressing a polypeptide in a C. elegans amphid sheath cell; and (4) expressing a polypeptide in a C. elegans containing the agent is useful for preventing or reducing composition containing the agent is useful for preventing or reducing composition containing the agent is useful for preventing the composition of plants, or for treating or reducing the likelihood of nematode sequence such as anaemia, malnutrition, pulmonary disorders, nutritional disorders, hyperinfection in immunocompromised individuals, nematocide, trichinosis, onchocerciasis (river blindness) and lymphatic filariasis. The VA genes are located on chromosome IV of the C. elegans genome. The present sequence is the VAP-2 protein

ents for treating, preventing or reducing nematode ants or individuals. ģ transgenic nematode having a transgene that regulates the expression nematode secretory product, useful for screening anti-nematode ts for treating, preventing or reducing nematode infestation in

aim 73; Fig 1; 105pp; English.

The invention relates to a transgenic nematode, the cells of which contain a transgene comprising a regulatory element of a gene that encodes a nematode secretory product or its homologue operably linked to a DNA sequence encoding a detectable marker. Also included are (1) identifying a compound that inhibits a nematode secretion pathway, for use as an anti-nematode agent for use in preventing or reducing mematode infestation of a plant (the agent may be used to treat the soil or seed); (2) a vector comprising; (a) a vap-1 or vap-2 polynucleotide encoding wat least 10 consecutive residues of VAP-1/VAP-2 or at least 50 % identity to VAP-1 or VAP-2; (iii) has a sequence comprising the Caenorhabditis elegans vap-1 or vap-2 promoter operably linked to a polynucleotide encoding a detectable marker; or (b) a regulatory element comprising to the C. elegans vap family of genes operably linked to a polynucleotide encoding a detectable marker; (3) expressing a first polynucleotide in a C. elegans amphid sheath cell; and (4) expressing a polypeptide in a C. elegans amphid sheath cell; and (4) expressing a polypeptide in a C. elegans amphid sheath cell; and (4) expressing a polypeptide in a C. elegans amphid sheath cell; and (4) expressing a polypeptide in a C. elegans amphid sheath cell; and (4) expressing a polypeptide in a C. elegans amphid sheath cell; and (4) expressing a polypeptide in a C. elegans amphid sheath cell; and (4) expressing a polypeptide in a C. elegans amphid sheath cell; and (4) expressing a polypeptide in a C. elegans detectable marker; (3) expressing a polypeptide in a C. elegans detectable marker; (3) expressing a polypeptide in a C. elegans detectable marker; (3) expressing a polypeptide in a C. elegans detectable marker; (3) expressing a first polynucleotide in a c. elegans detectable marker; (3) expressing a polypeptide in a C. elegans detectable marker; (3) expressing a polypeptide in a C. elegans detectable marker; (3) expressing and (4) expressing a polypeptide in a C. elegans detectable marke

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int antigen;
ialaria; measle
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7-OCT-2001;
3-NOV-2001;
5-APR-2002;
7-OCT-2002;
                                                                                                                                                                                     tihelmintic; anti-HIV; antitubercular; tuberculostatic; antimalarial; rucide; tranquilizer; antibacterial; vaccine; recombinant antigen; okworm; immune response; HIV infection; tuberculosis; malaria; measlestanus; diphtheria; pertussis; polio; gene; ds.
                                                                                                                                                                                                                                                                    okworm antigen #14.
                                                                                                                                                                                                                                                                                                   -MAY-2005
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PSDB; ADY37253.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             15.2%; Score 191; DB 9;
Similarity 29.5%; Pred. No. 2.3e-10;
32; Conservative 31; Mismatches 83
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        EVA--AQQIAPCNDPLPINTSLAQNIARWLY----
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NEVIWOEGKACY----CDARPDSFCCDNLC 215
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; 2001US-0329533P.
; 2001US-0332007P.
; 2002US-0375404P.
; 2002WO-US033106.
                                                                         2004US-00825692
                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          -ANKFTDPRLYTFANLAYDKNTALGCHYAKCQGPDRIVISCMYNN-VVPD 183
                                                                                                                                                                                                                                                                                                                                                                                                                                      213
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       -FKDSEEETVLQQVSWYWVSASL 133
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 424;
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N-PSDB; AI
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Loukas
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(ASHC/
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Composition hookworm in
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                                            ADY37286
             5-MAY-2005
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               invention.
                                                                                                                                                                                                                                       124
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   sition useful for vaccinating or
orm in mammal, comprises copy of
fragments derived from hookworm,
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LOUKAS A.
WILLIAMSON
JONES B.
BETHONY J.
GOUD G.
BOTTAZZI M
MENDEZ S.
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ASHCOM J.
BDAMCHIAN
ZHAN B.
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ADY37255.
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                                                                                                                                                                                                                                                                                               LGPAQNMYKVDWDCNLEEVAAQQIAPCNDPLPINTSLAQNIARWLYFKDSEEET--VLQQ
                                                                                                                                                                                                                                       VSWYWVSASLGFMKGTKLDQFAN--QWAEPLANIANYRNRKVGCAHKICPAQQNMVVSCV
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Tones B,
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G, Bottazzi
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antigen or
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(LOUK/)
(JONE/)
(BETH/)
(BOTT/)
(MEND/)
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                                                                                                                                                                   The invention relates to a composition (I) comprising a copy of recombinant or synthetic antigen or their fragments derived from hookworm and a carrier. (I) is useful for vaccinating or eliciting an immune response against hookworm in a mammal, reducing blood loss in a patient infected with hookworm, reducing hookworm size or quantitative egg count or hookworm burden in a patient infected with hookworm, or decreasing L3 migration across skin of a mammal. The above method further involves chemically treating a hookworm-infected patient prior to the step of administering. (M1) is also useful for vaccinating patient against infectious diseases e.g. HIV, tuberculosis, malaria, measles, tetanus, diphtheria, pertussis or polio. This sequence corresponds to a protein of the invention.
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3-NOV-2001;
5-APR-2002;
7-OCT-2002;
5-SEP-2003;
                                                                                   Local
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BOTTAZZI M E.
MENDEZ S.
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LOUKAS A.
WILLIAMSON
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                                                                    13.2%;
Similarity 27.5%;
53; Conservative 2
                               VVVAVLLALFSYAEAGFCCPNSLSQSDSARQIFLDFHNDVRRNIALGNGLINWTVNADAV
                                                                                                                                          217 AA;
VPLLVLLAVSVDANSVRCGNNGM--TDEAROKFLDMHNGYRSOVAKGQAKDALSGNAP--
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ashcom J, E
Williamson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2001US-0329533P.
2001US-0332007P.
2002US-0375404P.
2002WO-US033106.
2003US-0505848P.
                                                                                                                                                                                                                                                                                                                                                                                                         SEQ ID NO 58;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2004US-00825692
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ָם
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Bdamchian M,
1 A, Jones B,
                                                                                                                                                                                                                                                                                                                                                                                                         227pp;
                                                                     28;
                                                                     Score 166; DB 9; 1
Pred. No. 3.4e-08;
3; Mismatches 106;
                                                                                                                                                                                                                                                                                                                                                                                                         English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Zhan B, Wang
Bethony J, (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Goud
                                                                                                          Length 217;
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                                                                           Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Hawdon J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Bottazzi
                                                                       32;
                                                                           Gaps
                                         64
        58
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                                PCR primers All2/301 and All2/302 were designed to be used to amplify the DNA encoding the 45 kD protein. In the reaction containing primer C All2/301 a unique band of approximately 650 bp was observed. The C approximately 650 bp band was not seen when primer All2/302 was used. The C cloned was grown and purified for use as a hybridisation probe to screen the pBTA879 cDNA library. pBTA879 is the sequence of one of these c clones. There is no methionine initation codon present in this region of the sequence so this clone probably does not represent the complete coding region. pBTA879 probably encodes a glycoprotein which is related to, but not identical to, the native glycoprotein isolated from H. C contortus. In order to isolate a cDNA clone coding for the full length C isolated from pBTA879. AAQ28094 is the sequence of one of these clones, PBTA 963. Once again, this clone does not contain an initiation c management of the complete on the field.) (Updated on 25-MAR-2003 to correct PN field.) (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                            Vaccines comprise antigens derived passive immunisation against roundy threadworm and hookworm.
                                                                                                                                                                                                                                                                                                                                                                                                                                20-AUG-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO9213889-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Haemonchus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 25-MAR-2003
26-JAN-1993
                                                                                                                                                                                                                                                                                                                                                                                                         -FEB-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              114
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                                                                                                                                                                                                                      11;
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                                                                                                                                                                                                                                                                                                                                                                                    1991;
                                                                                                                                                                                                                                                                                         2-299990/36
AAQ28094.
                                                                                                                                                                                                                                                                                                                                                  BIOTECH AUSTRALIA COMMONWEALTH SCI
                            to correct
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     vaccine;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        VSCVYGSPKLAP-
                                                                                                                                                                                                                      Fig
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                                                                                                                                                                                                                                                                                                                           Wagland BM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                contortus
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(first en
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                                                                                                                                                                                                                                                                                                                                                                                    91AU-00004486
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Location/Qualifier
12. .31
/note= "homologous
239. .256
/note= "see above"
                                                                                                                                                                                                                                                                                                                                                                                                         92WO-AU000040
                                                                                                                                                                                                                     ; dd98
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    parasitic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   entry)
                                                                                                                                                                                                                     English
                            field.)
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& IND RES
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     nematode;
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                                                                                                                                                                                                                                                        roundworm,
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                                                                                                                                                                                                                                                      parasitic whipworm,
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filarial w
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                                                                                                                                                                                                                                                        worm,
                                                                 full length fragment
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                                                            clones,
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440

Composition useful for vaccinating or hookworm in mammal, comprises copy of their fragments derived from hookworm,

eliciting immune recombinant or sy and carrier.

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                                              Hotez P,
Loukas A,
Mendez S;
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23
26
17
26
                                MP
                                                                             (HOTE/)
(ASHC/)
(BDAM/)
(ZHAN/)
(ZHAN/)
(WANG/)
(HAWD/)
(LOUK/)
(LOUK/)
(MILL/)
(BOTT/)
(MEND/)
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3-NOV-2001;
5-APR-2002;
7-OCT-2002;
5-SEP-2003;
                                                                                                                                                                                                                                                                                                                                                                                            )6
)Y37296
                               I;
                                                                                                                                                                                                                                                                                                            tihelmintic; anti-HIV; antitubercular; trucide; tranquilizer; antibacterial; vacookworm; immune response; HIV infection;
                                                                                                                                                                                                                                         -APR-2004;
                                                                                                                                                                                                                                                                                                                                                                                                             14
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                                                                                                                                                                                                                                                         -FEB-2005
                                                                                                                                                                                                                                                                        2005042232-A1
                                                                                                                                                                                                                                                                                        cylostoma duodenal
                                                                                                                                                                                                                                                                                                       tanue;
                                                                                                                                                                                                                                                                                                                                              okworm antigen
                                                                                                                                                                                                                                                                                                                                                              -MAY-2005
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                                2005-194935/20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  182
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    118
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  7
                                                                                                                    HAWDON J.
LOUKAS A.
WILLIAMSON
                                                                                                                                                                  HOTEZ P.
ASHCOM J.
                                                                                     GOUD G.
BOTTAZZI
                                                                               MENDEZ
                                                                                                                                                   ZHAN
                                                                                                    BETHONY J.
                                                                                                            JONES B
                                                                                                                                                           BDAMCHIAN M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               13.0%;
Similarity 26.6%;
9; Conservative :
                                                                                                                                                                                                                                                                                                      diphtheria;
                                                                                                                                                                                                                                                                                                                                                                                             standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                VGYDCEVEANTAAYAKECKFEHDPPEQRNYWGONL--WMLGGTNYSK--TESAKLSVQAW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SYAEAGECCPNSLSQSDSARQIFLDFHNDVRRNIALGNGLINWTVNADAVILGPAQNMYK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   YWVSASLG-
                                                                                                                                                                                                                                                                                                                                                                                                                                   YNPAGNRINHY I YDIGDPĊTTD--BDWQCTGCTCSKDEALCI
                                                                                                                                                                                                                                                                                                                                                                                                                                                   YGSPKLAPNEVIWQEGKACVCDARPDSFCCDNLCDTRDAASV 223
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  YWELKMFGVPDENILTMEVFDRGVGHYTQ----VAWQSSDKIGCAVEWCPTM--TLVACE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                VDWDCNLEEVAAQQIAPC---NDPLPINTSLAQNIARWL----YFKDSEEETVLQQVSW 126
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SCLKAAFCPTSDNGMTDBIRQIFVDKHNEYRSIIAKGQ--
                                                       Ashcom J, 1
Williamson
                                                                                                                                                                                          2001US-0329533P.
2001US-0332007P.
2002US-0375404P.
2002WO-US033106.
2003US-0505848P.
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                                                                                                                                                                                                                                                                                                                                                             entry)
                                                      Bdamchian M, A, Jones B,
                                                                                                                                                                                                                                                                                                       pertussis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 -FMKGTKLDQFANQWABPLANIANYRNRKVGCAHKICPAQQNMVVSCV 181
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Pred. No. 1.7e-07; 24; Mismatches 109
                                                                                                                                                                                                                                                                                                                                                                                             216
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 163;
                                                                                                                                                                                                                                                                                                                                                                                              AA.
                                                                                                                                                                                                                                                                                                       polio.
                                                      Zhan B,
Bethony
                                                                                                                                                                                                                                                                                                             tuberculostatic; antimalari
accine; recombinant antigen;
tuberculosis; malaria; mea
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 2;
                                                     Wang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               109;
                                                       Goud
                                                            Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 440;
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                                                      Hawdon
G, Bot
                                                      lon J;
Botta
                                                                                                                                                                                                                                                                                                                              antimalarial;
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                                                      azzi
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                                                                                                                                                                                                                                                                                                               measle
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to a composition (I) comprising a copy of recombinant or synthetic antigen or their fragments derived from hookworm and a carrier. (I) is useful for vaccinating or eliciting an immune response against hookworm in a mammal, reducing blood loss in a patient infected with hookworm, reducing hookworm size or quantitative egg count or hookworm burden in a patient infected with hookworm, or decreasing L3 migration across skin of a mammal. The above method further involves chemically treating a hookworm-infected patient prior to the step of administering. (M1) is also useful for vaccinating patient against infectious diseases e.g. HIV, tuberculosis, malaria, measles, tetanus, for the step of the step o
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              administering. (M1) infectious diseases diphtheria, pertussi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  the
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                                17-OCT-2001;
23-NOV-2001;
26-APR-2002;
17-OCT-2002;
26-SEP-2003;
                                                                                                                                                                                                                                                                 virucide;
hookworm;
tetanus; d
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297
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                                                                                                                                                                                                                                                                                                                    antihelmintic;
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                                                                                                                                    94
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imilarity 27.4%;
; Conservative
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TELECOMMUNICATION INFORMATION:
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TELEFAX: (703) 683-4109
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CITY:
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                                                                                                                                                                                                                                                                                                                                                      COUNTRY: United ZIP: 22313-0299
                                                                                                                                                                                                                                                                                                 SOFTWARE:
                                                                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE:
                                                                                                                                                                                                                                                                              APPLICATION NUMBER:
                                                                                                                                                                                             FILING DATE:
                                                               TOPOLOGY:
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                                                                                    ENGTH:
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h 13.2%;
Similarity 26.6%;
59; Conservative
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: Virginia
RY: United S
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CHARACTERISTICS:
                                                    TYPE:
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IBM PC compatible
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INFORMATION:
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  re 165; DB 1;
d. No. 3e-09;
Mismatches 109;
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                                                                                                            US-08-
                                               Query Ma
Best Loc
Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY: United States of America ZIP: 20007-5109

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Ve
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                         APPLICATE: 06-FEB-108
FILING DATE: 06-FEB-109
ATTORNEY/AGENT INFORMATION:
NAME: Bent, Stephen A
REGISTRATION NUMBER: 29,768
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 672-5300
TELEPHONE: (202) 672-5399
                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US 07/930,686
FILING DATE: 06-OCT-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: AU PK4486
FILING DATE: 06-FEB-1991
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/AU92/00040
                                                                                                                                                                                    TELEX: 904136
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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CORRESPONDENCE ADDRESS:
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nce 12, Application US/08460998
t No. 5942413
                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US/08/460,998 FILING DATE: 05-JUN-1995 IOR APPLICATION DATA:
                                                            Local
                                                                                                               60-998-12
                                                                                                                             TOPOLOGY: 1
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                                                                Similarity
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                                                                                                                                         : 440 amino acids amino acid GY: linear
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SYAEAGFCCPNSLSQSDSARQIFLDFHNDVRRNIALGNGLINWTVNADAVILGPAQNMYK 74
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sharp, Phillip J
Wagland, Barry M
Cobon, Gary S
                                               13.2%; Score 165; DB 1; I larity 26.6%; Pred. No. 3e-09; Conservative 24; Mismatches 109;
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                                                                                MOLECULE TYPE 08-450-4828-125
                                                                                                                                                 INFORMATION
                                                                                                                                                                                                                                         FILING DATE: 10-NOV-1993
APPLICATION NUMBER: 08/060
FILING DATE: 11-MAY-1993
APPLICATION NUMBER: 07/996
FILING DATE: 24-DEC-1992
APPLICATION NUMBER: 07/881
FILING DATE: 11-MAY-1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL
                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION DATA
APPLICATION NUMBER: US
FILING DATE: 26-MAY-19
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08
PILING DATE: 23-DEC-19
APPLICATION NUMBER: 08
                                                                                                                                                                        TELECOMMUNICATION INFORMATION: TELEPHONE: (213) 489-1600 TELEFAX: (213) 955-0440
                                                                                                                                       SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NERAL INFORMATION:
APPLICANT: MATTHEW
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER
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CORRESPONDENCE ADDRE
ADDRESSEE: Lyon &
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDIUM TYPE: 81
                                                                                                                                                                                                        NAME: BIGGS, SUZANNE L. REGISTRATION NUMBER: 30 REFERENCE/DOCKET NUMBER:
                                                                                                   TOPOLOGY:
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                                                                                                                           LENGTH:
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CHARACTERISTICS:
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California
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IBM Compatible
SYSTEM: IBM P.C
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                                  10.9%; larity 24.9%; Conservative
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26-MAY-1995
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                                  Score 137; DB Pred. No. 1.3e 0; Mismatches
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                                            DB :
                                2; I
-06;
103;
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                                  Indels
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                                  30;
                                 Gaps
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APPLICANT
TITLE OF
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APPLICATION NUMBER: 07/99
FILING DATE: 24-DEC-1992
ATTORNEY/AGENT INFORMATION:
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APPLICATION NUMBER: US
FILING DATE: 10-NOVEMB
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MEDIUM TYPE: 3.5" Diskette,
MEDIUM TYPE: storage
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RIOR APPLICATION DATA:
APPLICATION NUMBER: 0:
FILING DATE: 11-MAY-1:
APPLICATION NUMBER: 0:
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                                                                                                                                                                               TOPOLOGY:
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RMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                    TELEFAX:
                                                                                                                                                                                                                                                                                                                 NAME: BIGGS, SUZANNE L.
REGISTRATION NUMBER: 30,158
REFERENCE/DOCKET NUMBER: 203/226
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OPERATING SYSTEM SOFTWARE: Word
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                                                                                                     Local Similarity
                                                                                                                                                                                                                                                                                     IECOMMUNICATION INFORMATION: TELEPHONE: (213) 489-1600
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EEVAAQQ-IAPCNDPLPINTSLAQNIARWLYFKDS---
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                                                         CCPNSLSQSDSARQIFLDFHNDVRRNIALGNGLINWTVNADAVILGPAQNMYKVDWDCNL 81
                             CKONNGSMTNELRRRFLRLHNGYRSILALGHVNISEESNETFLYAHRASRMRILDYDCDA 64
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11-MAY-1992
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11-MAY-1993
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24-DEC-1992
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10-NOVEMBER-1993
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                                                                                    Score 137; DB 2;
Pred. No. 1.3e-06;
30; Mismatches 103;
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RESULT 6
US-08-450-482
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APPLICATION NUMBER: 08/173
FILING DATE: 23-DEC-1993
APPLICATION NUMBER: 08/151
FILING DATE: 10-NOV-1993
APPLICATION NUMBER: 08/060
FILING DATE: 11-MAY-1993
APPLICATION NUMBER: 07/996
FILING DATE: 24-DEC-1992
APPLICATION NUMBER: 07/881
FILING DATE: 11-MAY-1992
APPLICATION NUMBER: 07/881
FILING DATE: 11-MAY-1992
APPLICATION NUMBER: 07/881
FILING DATE: 11-MAY-1992
ATTORNEY/AGENT INFORMATION:
NAME: BIGGS, SUZANNE L.
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                                                                                                                                                               TYPE:
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CHARACTERISTICS:
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Similarity 24.9%;
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Pred. No. 1.3e-06;
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ATTORNEY/AGENT INFORMATION:
NAME: BIGGS, SUZANNE L.
REGISTRATION NUMBER: 30,158
REFERENCE/DOCKET NUMBER: 203/
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEPHONE: (213) 955-0440
                                                                                                                                                                                                                                  TELEFAX: (213) 955-044
TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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CORRESPONDENCE ADDRESSEE: Lyon &
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                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US/08/151, US/FILING DATE: 10-NOVEMBER-1993
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/060, 433
FILING DATE: 11-MAY-1993
APPLICATION NUMBER: 07/881, 721
FILING DATE: 11-MAY-1992
APPLICATION NUMBER: 07/996, 972
APPLICATION NUMBER: 07/996, 972
FILING DATE: 24-DEC-1992
FILING DATE: 24-DEC-1992
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TOPOLOGY: LINFATOLOGY:
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MEDIUM TYPE: 3.5" Diskette,
MEDIUM TYPE: storage
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ice 125, Application US/08151064D
'No. 6962795
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Similarity 24.9%;
54; Conservative :
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California
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                       EEVAAQQ-IAPCNDPLPINTSLAQNIARWLYFKDS----EEETVLQQVSWYWVSASLGFM 136
                                                      | |: | :: |: || || | :|||: |: | : | :|:||: |CKQNNGSMTNELRRRFLRLHNGYRSILALGHVNISEESNETFLYAHRASRMRILDYDCDA 66
ĖGSAYESA I KQČSSNKSSSAEYDENV-
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Pred. No. 1.3e-0
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TELEPHONE: (21)
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                                                                                                                                                                                                                                             NAME: BIGGS, REGISTRATION
                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER:
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                                             CKQNNGSMTNELRRFLRLHNGYRSILALGHVNISEESNETFLYAHRASRMRILDYDCDA
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APPLICANT: MATTHEW MOTITLE OF INVENTION: INVE
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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REGISTRATION NUMBER: 30,1
REFERENCE/DOCKET NUMBER:
TELECOMMUNICATION INFORMATIC
TELEPHONE: (213) 489-1600
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COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Di
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APPLICATION NUMBER: US/(
FILING DATE: 10-NOVEMBER
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/(
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APPLICATION NUMBER: 07/8
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APPLICATION NUMBER: 07/9
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Pred. No. 1.4e-06;
); Mismatches 103
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VCDARPDSFCCDNLCD

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RESULT 10
US-08-450-482B-115
; Sequence 115, Application
; Patent No. 6818616
; GENERAL INFORMATION:
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COMPUTER READABLE
TYPE: 3.
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                                                                                                                                                                                                                                                                                          TELECOMMUNICATION
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COMPUTER: IBM Compatil
OPERATING SYSTEM: IBM
SOFTWARE: Word Perfectures
                                                                                                                                                                                                                                                                                                                                              FILING DATE: 10-NOV-1993
APPLICATION NUMBER: 08/060
FILING DATE: 11-MAY-1993
APPLICATION NUMBER: 07/996
FILING DATE: 24-DEC-1992
APPLICATION NUMBER: 07/881
FILING DATE: 11-MAY-1992
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STREET:
CITY: I
STATE:
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DLECULE TYPE:
450-482B-115
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                                                                                                                                             Local
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Similarity 24.8%;
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CHARACTERISTICS:
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26-MAY-1995
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11-MAY-1992
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Pred. No. 1.7e
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RESULT 11
US-08-151-064D-113
; Sequence 113, Application US/08151064D
; Patent No. 6962795
; GENERAL INFORMATION:
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COMPUTER READABLE FORM:

MEDIUM TYPE: 3.5" Diskette, 1.44

MEDIUM TYPE: storage

COMPUTER: IBM Compatible

COMPUTER: IBM Compatible

OPERATING SYSTEM:

SOFTWARE: Word Perfect 5.1

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/151,06,

FILING DATE: 10-NOVEMBER-1993

CLASSIFICATION DATA:

APPLICATION NUMBER: 08/060,433

FILING DATE: 11-MAY-1993

APPLICATION NUMBER: 07/881,721

FILING DATE: 11-MAY-1992

APPLICATION NUMBER: 07/996,972

FILING DATE: 24-DEC-1992

ATTORNEY/AGENT INFORMATION:

NAME: RIGGS, SUZANNE L.
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TITLE OF IN
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                                                                                                                                                                                                                                                                                                                      NAME: BIGGS, SUZANNE L.
REGISTRATION NUMBER: 30,158
REFERENCE/DOCKET NUMBER: 203,
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
                                                                                                                                                           Local
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STATE: California
COUNTRY: U.S.A.
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                                                                                                                                                                                                                                                   TYPE:
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TLE OF INVENTION: NOVEL NEUTR
MBER OF SEQUENCES: 130
RESPONDENCE ADDRESS:
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AMINO ACID
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| ITEESEDYDLYDLLYAPTASKMRYLEYDCEAEKSAYESAKKCQTTAFSSTKYDENLQVIE
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                                                                                                                 SAAVVVAVLLALFSYAEAGFCCPNSLSQ-----SDSARQIFLDFHNDVRRNIALGNGLIN 56
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633 West Fifth
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ilarity 24.8%;
Conservative 35
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10-NOVEMBER-1993
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Pred. No. 1.7e-06;
35; Mismatches 102;
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RESULT 12
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Best Local &
Matches 57
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INFORMATION FOR SEQ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                OPERATING SYSTEM:
SOFTWARE: Patent
CURRENT APPLICATION
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FILING DATE: 23-MAY-1996
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                             TOPOLOGY: 1 MOLECULE TYPE:
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TITLE OF INVENTION: PARASITIC
TITLE OF INVENTION: ANTIGEN 5
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CITY: I
STATE:
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                                                                                                                                                                                                                                                                                                              LENGTH:
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UTER: IBM PC compatible
ATING SYSTEM: PC-DOS/MS-I
WARE: PatentIn Release #1
T APPLICATION DATA:
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Similarity 23.5%;
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ID NO:
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Mismatches 91;
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TELEFAX: (303) 863-0223 INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
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NAME: Connell, Gary J.
REGISTRATION NUMBER: 32,0
REFERENCE/DOCKET NUMBER:
TELECOMMUNICATION INFORMATION

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APPLICANT:
TITLE OF INV
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APPLICATION NUMBER: PCT/US96/077
FILING DATE: 23-MAY-1996
CLASSIFICATION:
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CORRESPONDENCE ADDRESS:
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96-07709-25
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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STREET: 1700 Lincoln
CITY: Denver
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VENTION: PARASITIC HELMINTH VENOM
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TOPOLOGY: linear
MOLECULE TYPE: pr
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COMPUTER READABLE
MEDIUM TYPE: Fl
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/
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CORRESPONDENCE ADDRESS:
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CLASSIFICATION: 514
PRIOR APPLICATION DATA:
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SOFTWARE: PatentIn Release
CURRENT APPLICATION DATA:
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (703) 836-9300
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ATTORNEY/AGENT
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 e 10, Application US/084
No. 5942413
L INFORMATION:
ICANT: Sharp, Phillip J
ICANT: Wagland, Barry N
ICANT: Cobon, Gary S
ICANT: Cobon, Gary S
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T: Cobon, Gary S
INVENTION: Nema
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Similarity 24.4%;
64; Conservative
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United States
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INFORMATION:
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Ve
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/460,998
FILING DATE: 05-JUN-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/930,686
FILING DATE: 06-OCT-1992
PRIOR APPLICATION DATA:
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TOPOLOGY: linear
MOLECULE TYPE: pro-
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FILING DATE: 06-FEB-1991
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/AU92/00
FILING DATE: 06-FEB-1992
ATTORNEY/AGENT INFORMATION:
NAME: Bent, Stephen A
REGISTRATION NUMBER: 29,768
REGISTRATION INFORMATION:
TELEPHONE: (202) 672-5399
TELEPHONE: (202) 672-5399
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QUENCE CHARACTERISTICS:
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; Publication No. US20040052817A1
; GENERAL INFORMATION:
APPLICANT: Universiteit Gent
; TITLE OF INVENTION: Ostertagia vaccine
; FILE REFERENCE: UniGent Ostertagia
; CURRENT APPLICATION NUMBER: US/10/243,319C
; CURRENT FILING DATE: 2002-09-13
; NUMBER OF SEQ ID NOS: 14
; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 10
; LENGTH: 231
; TYPE: PRT
; ORGANISM: Ostertagia ostertagi
US-10-243-319C-10
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          ICANT: Universiteit Gent
E OF INVENTION: Ostertagia vaccine
REFERENCE: UniGent Ostertagia
ENT APPLICATION NUMBER: US/10/243,319C
ENT FILING DATE: 2002-09-13
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Result No.

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e 50, Ap	nce 50. Ap	e 50, App	0, Ap	e 50, Ap	e 50, Ap	e 50, Ap	e 50, Ap	e 50,	e 50,	e 50, A	e 50, A			Sequence 5, Appli	, Ap	Sequence 47, Appl	Sequence 2, Appli

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Watch 100.0%;
Local Similarity 100.0%;
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nce 8, Application US/10243319C cation No. US20040052817A1 AL INFORMATION:
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Pred. No. 5.6e-119;
Mismatches 0;
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APPLICANT: Mendez, Susana
TITLE OF INVENTION: Hookworm Vaccine
FILE REFERENCE: 03740007aa
CURRENT APPLICATION NUMBER: US/10/825,692
CURRENT FILING DATE: 2004-04-16
PRIOR APPLICATION NUMBER: US 60/329,533
PRIOR APPLICATION NUMBER: US 60/329,533
PRIOR APPLICATION NUMBER: US 60/332,007
PRIOR APPLICATION NUMBER: US 60/332,007
PRIOR APPLICATION NUMBER: US 60/375,404
PRIOR FILING DATE: 2002-04-26
PRIOR FILING DATE: 2002-04-26
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LENGTH: 236
TYPE: PRT
ORGANISM: Ostertagia ost
FEATURE:
NAME/KEY: misc_feature
LOCATION: (37)..(37)
OTHER INFORMATION: The
FEATURE:
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           SEQ ID NO 18
LENGTH: 42
TYPE: PRT
ORGANISM:
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                                                               PRIOR FILING DATE: 2002-04-26
PRIOR APPLICATION NUMBER: PCT
PRIOR FILING DATE: 2002-10-17
NUMBER OF SEQ ID NOS: 114
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                                                       SOFTWARE:
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GANISM: Ancylostoma
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Local Similarity 52.5%;
hes 125; Conservative
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Loukas, Alex
Williamson,
Jones, Briar
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Bdamchian, Ma
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Wang, Yan
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Goud, Gaddam
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APPLICANT: Goud, Gaddam
APPLICANT: Botazzi, Maria E.
APPLICANT: Mendez, Susana
TITLE OF INVENTION: Hookworm Vaccine
FILE REFERENCE: 03740007aa
CURRENT APPLICATION NUMBER: US/10/825,692
CURRENT FILING DATE: 2004-04-16
PRIOR APPLICATION NUMBER: US 60/329,533
PRIOR FILING DATE: 2001-10-17
PRIOR APPLICATION NUMBER: US 60/332,007
PRIOR APPLICATION NUMBER: US 60/332,007
PRIOR APPLICATION NUMBER: US 60/332,007
PRIOR APPLICATION NUMBER: US 60/375,404
PRIOR APPLICATION NUMBER: PCT US02/33106
PRIOR APPLICATION NUMBER: PCT US02/33106
PRIOR FILING DATE: 2002-10-17
NUMBER OF SEQ ID NOS: 114
SOFTWARE: PatentIn version 3.2
SEQ ID NO 67
LENGTH: 425
TYPE: PRT
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US-10-825-692-67
; Sequence 67, Applicat
; Publication No. US200
; GENERAL INFORMATION:
; GENERAL TOTAL Hotez, Pe
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Similarity 30.9%; Pred. No. 2.6e-19;
76; Conservative 35; Mismatches 97;
                                                                                             21.9%; Score 274; DB 5; l
Similarity 30.3%; Pred. No. 5.3e-19;
74; Conservative 37; Mismatches 97;
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GLINWTVNADAVILGPAQNMYKVDWDCNLEEVAAQQIAPCNDPLPINTSLAQNIARWLY-
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Loukas, Alexander
Williamson, Angela
Jones, Brian
                                                       SAAVVVAVLLAL----FSYAEAGFCCPNSLSQSDSARQIFLDFHNDVRRNIALG----N
                               Application US/10825692 vo. US20050042232A1
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Wang, Yan
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Bdamchian, Mahnaz
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RESULT 6
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                                                                                        Sequence 2, Application US/10825692 Publication No. US20050042232A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER:
CURRENT FILING DATE: 2003-(
NUMBER OF SEQ ID NOS: 8
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 5
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Publication No. US20030126625A1
GENERAL INFORMATION:
APPLICANT: Liu, et al.
TITLE OF INVENTION: Screens and Assays for
TITLE OF INVENTION: Parasitic Nematodes
FILE REFERENCE: 2002630-0012
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ORGANISM: Artificial
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                                                                                                                                                                                                                                                DSEESTVLQQVSWYWVSASLGFMKGTKLDQ-
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                                   Ashcom, Jan
Bdamchian,
Zhan, Bin
Wang, Yan
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Hawdon, John
Loukas, Alexander
Williamson, Angel
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larity 31.3%;
Conservative
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Pred.
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LENGTH: 424
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: Hookworm FILE REFERENCE: 03740007aa CURRENT APPLICATION NUMBER: UCURRENT FILING DATE: 2004-04 PRIOR APPLICATION NUMBER: US PRIOR FILING DATE: 2001-10-17
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Best I
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OR APPLICATION NUMBER: US 60/329,533

OR FILING DATE: 2001-10-17

OR APPLICATION NUMBER: US 60/332,007

OR FILING DATE: 2001-11-23

OR APPLICATION NUMBER: US 60/375,404

OR APPLICATION NUMBER: PCT US02/33106

OR APPLICATION NUMBER: PCT US02/33106

OR FILING DATE: 2002-10-17

SER OF SEQ ID NOS::114
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OF INVENTION: Hookworm Vaccine
REFERENCE: 03740007aa
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                                QNMVVSCVYGSPKLAPNEVIWQEGKACV----CDARPDSFCCDNLCDT-RDAASVRHQC
                                                                DPSVKIEPTLSGWWSGAK---KNGVGPDNKYTGGGLFAFSNMVYSETTKLGCAYKVCGTK
                                                                                              EETVLQQ--VSWYWVSASLGFMKGTKLD-QFANQWAEPLANIANYRNRKVGCAHKICPAQ 173
                                                                                                                                                               WIVNADAVILGPAQNMYKVDWDCNLEEVAAQQIAPCNDPLPINTSLAQNIARWLYFKDSE 116
                                                                                                                                                                                             MFSPVVVSVVFTIAFCNASPARDSFGCSNS-GITDSDRQAFLDFHNNARRRVAKG-----
                                                                                                                                                                                                                             MSAAVVVAVLLAL----FSYAEAGFCCPNSLSQSDSARQIFLDFHNDVRRNIALGNGLIN 56
--LAVSCIYNGVGYITNQPMWETGQACQTGADCSTYKNSGCEDGLCTKGPDVPETNQQC
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Bethony, Jeft:
Gaddam
                                                                                                                            -LEDSNSGKLNPAKNMYKLSWDCAMEQQLQDAIQSCPSGFAGIQGVAQNTMSWSSSGGYP
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illarity 30.5%;
Conservative 36
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entIn version 3.2
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Pred. No. 6.9e-18;
5; Mismatches 106
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  227
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RESULT US-10-8 APPLICANT: Mendez, Susana
TITLE OF INVENTION: Hookworm Vaccine
FILE REFERENCE: 03740007aa
CURRENT APPLICATION NUMBER: US/10/825,692
CURRENT FILING DATE: 2004-04-16
PRIOR APPLICATION NUMBER: US 60/329,533
PRIOR FILING DATE: 2001-10-17
PRIOR APPLICATION NUMBER: US 60/332,007
PRIOR FILING DATE: 2001-11-23
PRIOR APPLICATION NUMBER: US 60/375,404 Sequer Public GENERA APPLICANT:
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APPLICANT: APPLICANT: APPLICANT: APPLICANT: APPLICANT: APPLICANT: $\boldsymbol{\omega}$ ence 56, Applica ication No. US20 RAL INFORMATION: 25-692-56 Hotez, Peter Bethony, Jerr Goud, Gaddam Zhan, Bin Wang, Yan Hawdon, John Loukas, Alexander Application US/10825692 Botazzi, Jones, Brian Williamson, Ashcom, James Bdamchian, Mahnaz Maria B. Jeffrey Angela US/10/825,692

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RESULT 8
US-10-051-
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APPLICANT: Liu, et al.
TITLE OF INVENTION: Screens and Assay:
TITLE OF INVENTION: Parasitic Nemator
FILE REFERENCE: 2002630-0012
CURRENT APPLICATION NUMBER: US/10/051
NUMBER OF SEQ ID NOS: 8
SOFTWARE: PatentIn Ver. 2.1
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PRIOR APPLICATION NUMBER: PCT U
PRIOR FILING DATE: 2002-10-17
NUMBER OF SEQ ID NOS: 114
SOFTWARE: Patentin version 3.2
SEQ ID NO 56
LENGTH: 422
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Matches 76
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Matches 62
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FEATURE:
OTHER INFORMATION: Des
OTHER INFORMATION: Act
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ORGANISM: Ancylostoma
.0-825-692-56
                                                                                                                                                                                         OTHER INFORMATION:
OTHER INFORMATION:
.0-051-644B-3
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                                                                      WDCNLEEVAAQQIAPCND---PLPINTSLAQNIARWLYFKDSEEETV-LQQVSWYWVSAS
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                         LGFMKGTKLDQ-
                                             WDCELASLAANWSTSCPQHFMPQSVLGSNAQLFKRFYFYFDGHDSTVHMRNAMKYWWQQ-
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CNKSTITQLQ--QEIILTTHNELRRSLAFGKQRNKRGLMN-
-GEEKGNE-DQKNRFYARRNYFGWANMAKGKTYRVGCSYIMCGDGESALFTCLYNEKAQC
                                                                                                                     CCPNSLSQSDSARQIFLDFHNDVRRNIALG--
                                                                                                                                           17.3%; larity 29.1%; Conservative
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                        - FANQWAEPLANIANYRNRKVGCAHKI CPAQQNMVVSCVYGSPKLA
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e-13;
77;
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GENERAL INFORMALLON:
APPLICANT: Liu, et al.
TITLE OF INVENTION: Screens and Assays for Agritle of INVENTION: Parasitic Nematodes
FILE REFERENCE: 2002630-0012
CURRENT APPLICATION NUMBER: US/10/051,644B
UMBER OF SEQ ID NOS: 8
SOFTWARE: PatentIn Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; FEATURI
; OTHER :
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US-10-051-
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Matches
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                                                                                                                                                                         Sequence 26, Applica Publication No. US20 GENERAL INFORMATION:
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APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT: Mendez, Susana
TITLE OF INVENTION: Hookworm Vaccine
FILE REFERENCE: 03740007aa
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Similarity 26.5%;
55; Conservative 3
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Bethony, Jeff:
Tand, Gaddam
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Wang, Yan
Hawdon, John
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Williamson, Angela
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                                                                                                                                        Ashcom, Jam
Bdamchian,
                                Botazzi,
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                                                           Jeffrey
                                                                                                                                            Mahnaz
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Pred. No. 1.4e-12;
36; Mismatches 76;
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US-10-825-692-28
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Best Local
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ORGANISM: F
-10-825-692-2
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                               NUMBER OF
                                           PRIOR
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CURRENT APPLICATION NUMBER:
CURRENT FILING DATE: 2004-
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           ID NO
                                    RENT FILING DATE: 2004-04-16

OR APPLICATION NUMBER: US 60/329,533

OR FILING DATE: 2001-10-17

OR APPLICATION NUMBER: US 60/332,007

OR FILING DATE: 2001-11-23

OR APPLICATION NUMBER: US 60/375,404

OR FILING DATE: 2002-04-26

OR APPLICATION NUMBER: PCT US02/33106

OR FILING DATE: 2002-10-17
                                                                                                                                                           CANT: Botazzi, Maria E.

CANT: Mendez, Susana

OF INVENTION: Hookworm Vaccine
                               FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US 60/375,404 FILING DATE: 2002-04-26 APPLICATION NUMBER: PCT US02/3310 FILING DATE: 2002-10-17
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Similarity 29.5%;
62; Conservative
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Hawdon, John
Loukas, Alexander
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Bethony,
Goud, Gao
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lo. US20050042232A1
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; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 58
; LENGTH: 217
; TYPE: PRT
; ORGANISM: Ancylostoma ceylanicum
US-10-825-692-58
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                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/10/825,692
CURRENT FILING DATE: 2004-04-16
PRIOR APPLICATION NUMBER: US 60/329,533
PRIOR FILING DATE: 2001-10-17
PRIOR FILING DATE: 2001-11-23
PRIOR APPLICATION NUMBER: US 60/375,404
PRIOR FILING DATE: 2002-04-26
PRIOR APPLICATION NUMBER: PCT US02/33106
PRIOR FILING DATE: 2002-10-17
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TITLE OF INVENTION: Hookworm Vaccine
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Similarity 27.5%;
3; Conservative 2
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ILGPAQNMYKVDWDCNLEEVAAQQIAPCNDPLPINTSLAQNIARWL-YFKDSEBETVLQQ
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                                                         VVVAVLLALFSYAEAGFCCPNSLSQSDSARQIFLDFHNDVRRNIALGNGLINWTVNADAV 64
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Bdamchian,
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No. US20050042232A1
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Loukas, Alexander
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Pred. No. 2.3e
28; Mismatches
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PRIOR FILING DATE: 2001-10-17
PRIOR APPLICATION NUMBER: US 60/332,007
PRIOR FILING DATE: 2001-11-23
PRIOR APPLICATION NUMBER: US 60/375,404
PRIOR FILING DATE: 2002-04-26
PRIOR APPLICATION NUMBER: PCT US02/33106
PRIOR FILING DATE: 2002-10-17
NUMBER OF SEQ ID NOS: 114
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Best Local &
Matches 68
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CURRENT FILING DATE: 2004-04-16
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0-825-692-
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CANT: Botazzi, Maria E.
CANT: Mendez, Susana
OF INVENTION: Hookworm Vaccine
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Loukas, Alexander
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ilarity 27.4%;
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GENERAL INFORMATION: APPLICANT: Hotez, P

APPLICANT:
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APPLICANT:

Hotez, Percana Ashcom, James Bdamchian, Mah

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LICANT: LICANT:

Hawdon, John Loukas, Alexander Williamson, Angel

Sequence 20, Appublication No.

Application US/10825692 No. US20050042232A1

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APPLICANT: Mendez, Susana
TITLE OF INVENTION: Hookworm Vaccine
FILE REFERENCE: 03740007aa
CURRENT APPLICATION NUMBER: US/10/825,692
CURRENT FILING DATE: 2004-04-16
PRIOR APPLICATION NUMBER: US 60/329,533
PRIOR FILING DATE: 2001-10-17
PRIOR APPLICATION NUMBER: US 60/332,007
PRIOR APPLICATION NUMBER: US 60/375,404
PRIOR APPLICATION NUMBER: US 60/375,404
PRIOR APPLICATION NUMBER: PCT US02/33106
PRIOR FILING DATE: 2002-04-26
PRIOR APPLICATION NUMBER: PCT US02/33106
PRIOR FILING DATE: 2002-10-17
NUMBER OF SEQ ID NOS: 114
SOFTWARE: PatentIn version 3.2
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ESULT 15
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Sequence 20, Ap
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Hawdon, John
Loukas, Alexander
Williamson, Angela
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                                                             VEAC---SIMCYVVCQYGPAGNMMGKDIYEKGEPC-
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Wang, Yan
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Bdamchian, Mahnaz
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Pred. No. 3.5e-07;
35; Mismatches 84;
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APPLICANT: Jones, Brian
APPLICANT: Bethony, Jeffrey
APPLICANT: Goud, Gaddam
APPLICANT: Botazzi, Maria E.
APPLICANT: Mendez, Susana
TITLE OF INVENTION: Hookworm Vaccine
FILE REFERENCE: 03740007aa
CURRENT APPLICATION NUMBER: US/10/825,692
CURRENT FILING DATE: 2004-04-16
PRIOR APPLICATION NUMBER: US 60/329,533
PRIOR FILING DATE: 2001-10-17
PRIOR APPLICATION NUMBER: US 60/332,007
PRIOR APPLICATION NUMBER: US 60/332,007
PRIOR APPLICATION NUMBER: US 60/332,007
PRIOR APPLICATION NUMBER: US 60/375,404
PRIOR APPLICATION NUMBER: PCT US02/33106
PRIOR FILING DATE: 2002-04-26
PRIOR APPLICATION NUMBER: PCT US02/33106
PRIOR FILING DATE: 2002-10-17
NUMBER OF SEQ ID NOS: 114
SOFTWARE: Patentin version 3.2
SEQ ID NO 20
LENGTH: 218
TYPE: PRT
ORGANISM: Ancylostoma caninum
US-10-825-692-20
Search completed: March Job time: 166 secs
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/cgn2_6/ptodata/1/pubpaa/US09_NEW_PUB.pep:*
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/cgn2_6/ptodata/1/pubpaa/US11_NEW_PUB.pep:*
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IS-10-527-500-23
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RESULT 2
US-10-527-771-2
; Sequence 2, Application US/10527771
; Publication No. US20050271683A1
; GENERAL INFORMATION:
; APPLICANT: University Gent
; TITLE OF INVENTION: Ostertagia vac
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Sequence 59, Appl	Sequence 5200, Ap	Sequence 1660, Ap	Sequence 141, App	ce 5684,	ce 82,	ce 80,	ce 96,	e 84, App	e 796	e 7,	Sequence 8, Appli	equence 9, Ap	equence	ce 9, App	0	ce 65, Ap	ce 104, A	Sequence 72, Appl	

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RESULT 1 US-10-527-771-10 Sequence 10, Application US/10527771 Publication No. US20050271683A1 GENERAL INFORMATION: APPLICANT: University Gent TITLE OF INVENTION: Ostertagia vaccine FILE REFERENCE: 2002-015 CURRENT APPLICATION NUMBER: US/10/527,771 CURRENT FILING DATE: 2005-03-11 PRIOR APPLICATION NUMBER: US 10/243,319 PRIOR FILING DATE: 2002-09-13 NUMBER OF SEQ ID NOS: 27 SOFTWARE: PatentIn version 3.2 SEQ ID NO 10 LENGTH: 231 TYPE: PRT ORGANISM: Ostertagia ostertagi US-10-527-771-10 ches Match 100.0%; Score 1253; DB 6; Local Similarity 100.0%; Pred. No. 3.8e-123; les 231; Conservative 0; Mismatches 0; 181 181 121 121 61 13 \vdash MSAAVVVAVLLALFSYABAGFCCPNSLSQSDSARQIFLDFHNDVRRNIALGNGLINWTVN 60

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TITLE OF INVENTION: Adjuvancy and Immu
TITLE OF INVENTION: of Onchocerca Vol
FILE REFERENCE: 454-37
CURRENT APPLICATION NUMBER: US/11/153,
CURRENT FILING DATE: 2005-06-15
PRIOR APPLICATION NUMBER: 60/580,254
PRIOR FILING DATE: 2004-06-15
NUMBER OF SEQ ID NOS: 3
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ORGANISM: Ostertagia ost
FEATURE:
NAME/KEY: misc feature
LOCATION: (37) . (37)
OTHER INFORMATION: The 1
FEATURE:
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CURRENT APPLICATION NUMBER: US/10/527,7'
CURRENT FILING DATE: 2005-03-11
PRIOR APPLICATION NUMBER: US 10/243,319
PRIOR FILING DATE: 2002-09-13
NUMBER OF SEQ ID NOS: 27
SOFTWARE: Patentin version 3.2
                                                                                                                                                           SEQ ID NO
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TYPE: PRT
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RESULT 5
US-10-515-868-5
; Sequence 5, Application US/10515868
; Publication No. US20050282729A1
; GENERAL INFORMATION:
; APPLICANT: Hamilton, David W
; APPLICANT: Roberts, Kenneth P
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US-11-15
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US-11-153-222A-2
; Sequence 2, Application US/11153222A
; Publication No. US20060039921A1
; GENERAL INFORMATION:
; APPLICANT: Lustigman, Sara
; APPLICANT: MacDonald, Angus J.
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SEQ II
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TITLE OF INVENTION: Adjuvancy and Immune Potentiating
TITLE OF INVENTION: of Onchocerca Volvulus
FILE REFERENCE: 454-37
CURRENT APPLICATION NUMBER: US/11/153,222A
CURRENT FILING DATE: 2005-06-15
PRIOR APPLICATION NUMBER: 60/580,254
PRIOR FILING DATE: 2004-06-15
NUMBER OF SEQ ID NOS: 3
SOFTWARE: PatentIn version 3.3
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OF INVENTION:
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larity 22.1%;
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               CONTRACEPTIVES
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CURRENT APPLICATION NUMBER: US/11/153,22
CURRENT FILING DATE: 2005-06-15
PRIOR APPLICATION NUMBER: 60/580,254
PRIOR FILING DATE: 2004-06-15
NUMBER OF SEQ ID NOT
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LENGTH: 220
TYPE: PRT
ORGANISM: Onchocerca
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Publication No. US20060039921A1
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11-153-222A-3
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CURRENT APPLICATION NUMBER: US/10/515,869
CURRENT FILING DATE: 2004-11-24
PRIOR APPLICATION NUMBER: 60/383,628
PRIOR PILING DATE: 2002-05-28
PRIOR APPLICATION NUMBER: PCT/US03/16669
PRIOR PILING DATE: 2003-05-28
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0-515-868-5
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Pred. No. 6.8e-05;
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CURRENT FILING DATE: 2005-03-25
PRIOR APPLICATION NUMBER: 60/556,344
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TITLE OF INVENTION: Immortalized Human Tuberous
TITLS OF INVENTION: Angiomyolipoma Cell and Met
PILE REFERENCE: 24318-502
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ENT APPLICATION NUMBER: US/10/51
ENT FILING DATE: 2004-11-24
R APPLICATION NUMBER: 60/383,628
R FILING DATE: 2002-05-28
R FILING DATE: 2002-05-28
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ANISM: HOMOSAPIEN
15-868-6
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E OF INVENTION: CRISP POLYPEP
E OF INVENTION: CAPACITATION
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                                                                                 APPLICATION NUMBER: PCT
FILING DATE: 2003-05-28
R OF SEQ ID NOS: 14
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                                                                                                                                                                                                                                                                          NUMBER
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TYPE: PRT
ORGANISM:
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DR FILING DATE: 2002-07-18
DR APPLICATION NUMBER: US 10/052,586
DR FILING DATE: 2002-01-15
DR APPLICATION NUMBER: PCT/US01/06520
DR APPLICATION NUMBER: US 09/380,138
DR APPLICATION NUMBER: US 09/380,138
DR FILING DATE: 1999-08-25
DR APPLICATION NUMBER: PCT/US99/05028
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APPLICATION NUMBER:
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Godowski, Paul
Wood, William
Smith, Victoria
                                                                        APC---NDPLPINTSLAONI-ARWLYFKDSEEETVLQQVSWYWVSASLGFMKGTKLDQFA
                                                                                                                         SDSARQIFLDFHNDVRRNIALGNGLINWTVNADAVILGPAQNMYKVDWDCNLEEVAAQQI
                        NQWAEPLANIANYR-
                                                 ESCLWEHGPASLLPSIGONLGAHWGRYR-
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YPYEHECNPYCPFRCSGPVCTHYTQVVWATSNRIGCAINLC-
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Pred. No. 0.0011;
5; Mismatches 99;
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RESULT 11
US-10-506-443A-41
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TITLE OF INVENTION: CRISP POLYPEPTIDES AS (
TITLE OF INVENTION: CAPACITATION
FILE REFERENCE: 110.01860101
CURRENT APPLICATION NUMBER: US/10/515,868
CURRENT FILING DATE: 2004-11-24
PRIOR APPLICATION NUMBER: 60/383,628
PRIOR FILING DATE: 2002-05-28
PRIOR FILING DATE: 2003-05-28
PRIOR FILING DATE: 2003-05-28
                                     US-10-5
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SEQ ID NO 4
LENGTH: 2
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SOFTWARE: Pat
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Query
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SOFTWARE: PatentIn version 3.2
                                                                                                                      APPLICANT: Koslowski Dr., Michael TITLE OF INVENTION: Genetic Products DifferilE REFERENCE: 342-3PCT CURRENT APPLICATION NUMBER: US/10/506,443A CURRENT FILING DATE: 2004-09-01
                                                                                                                                                                                  APPLICANT: Sahin APPLICANT: Turec
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ORGANISM: HOMOSAPIEN
LO-515-868-4
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ORGANISM: Homo
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[CANT: Tureci Dr., Ozlem
[CANT: Koslowski Dr., Michael
] OF INVENTION: Genetic Products
REFERENCE: 342-3PCT
            Match
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  Similarity
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TYPE: PRT
ORGANISM:
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R APPLICATION NUMBER: US 6(
R FILING DATE: 2002-11-12
R APPLICATION NUMBER: US 6(
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Fischer, Laurent Berna...
Audonnet, Jean-Cristophe
Milward, Francis William
MIWARD, P. ARIASI POLYPEPTIDES A
MOTHODS OF USE
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                                                                PLPINTS---LAQNIARWLYFKD-SEEETVLQQVSWYWVS--
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Ribeiro, Jo
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US-10-4

53-372-102

Sequence 102, Applic Publication No. US20 GENERAL INFORMATION:

Application US/10453372 o. US20060003323A1

TITLE I

21402-589

THERAPEUTIC

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APPLICANT: Alsobrook, TITLE OF INVENTION: T

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RESULT 13
US-10-453-372-100
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SEQ ID NO 100
LENGTH: 497
TYPE: PRT
ORGANISM: Homo of 10-453-372-100
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APPLICATION NUMBER: 09/8:
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APPLICATION NUMBER:
FILING DATE: 2000-03
APPLICATION NUMBER:
FILING DATE: 2001-03
APPLICATION NUMBER:
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ing Prior Application data
! OF SEQ ID NOS: 1609
RE: CuraSeqList version 0.1
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APPLICATION NUMBER: 09/863776
FILING DATE: 2001-05-23
APPLICATION NUMBER: 60/208263
FILING DATE: 2000-05-31
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Similarity 21.0%;
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o. US20060003323A1
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RESULT 15
US-10-453-372-98
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                              FILE REFERENCE: 21402-589 A
CURRENT APPLICATION NUMBER: US/10/453
CURRENT FILING DATE: 2003-06-03
PRIOR APPLICATION NUMBER: 09/789390
PRIOR APPLICATION NUMBER: 60/185967
PRIOR FILING DATE: 2000-03-01
PRIOR APPLICATION NUMBER: 09/823187
PRIOR APPLICATION NUMBER: 60/195792
PRIOR APPLICATION NUMBER: 60/195792
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10-453-372-1
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R APPLICATION NUMBER: 60/195792
R FILING DATE: 2000-03-10
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R APPLICATION NUMBER: 60/199476
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RAPPLICATION NUMBER: 09,
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TH: 497
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APPLICATION NUMBER: 0
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APPLICATION NUMBER: 0 PILING DATE: 2001-03-APPLICATION NUMBER: 6
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R APPLICATION NUMBER: 09/863776

PREFILING DATE: 2001-05-23

PREFILING DATE: 2000-05-31

PREFILING DATE: 2000-05-31

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protein F11C7.3 [imported] - Caenorhabditis elegans
C;Species: Caenorhabditis elegans
C;Date: 10-May-2001 #sequence_revision 10-May-2001 #text_change
C;Accession: C89753
R;anonymous, The C. elegans Sequencing Consortium.
Science 282, 2012-2018, 1998
A;Title: Genome sequence of the nematode C. elegans: a platform
A;Reference number: A75000; MUID:99069613; PMID:9851916
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       tical protein T05A10.4 - Caenorhabditis elegans
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R;Sulston, J.
submitted to the EMBL Data Library, |
A;Reference number: Z19898
A;Accession: T24494
A;Status: preliminary; translated from type: DNA
A;Molecule type: DNA
A;Residues: 1-221 <WIL>
A;Cross-references: UNIPROT: P90959;
A;Experimental source: clone T05A10
C;Genetics:
A;Gene: CESP:T05A10.4
A;Map position: X
A;Introns: 18/3; 54/1; 106/3; 142/2;
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hypothetical protein ZK384.1 - C; Species: Caenorhabditis elega C; Date: 15-Oct-1999 #sequence_r C; Accession: T27833
R; Ainscough, R.
submitted to the EMBL Data Libr A; Reference number: Z20427
A; Reference number: Z20427
A; Accession: T27833
A; Status: preliminary; translat A; Molecule type: DNA
A; Residues: 1-196 <WIL>A; Cross-references: UNIPROT:062
A; Experimental source: clone ZK C; Genetics:
A; Gene: CESP: ZK384.1
A; Map position: 5
A; Introns: 58/3; 95/3; 157/2
C; Superfamily: yellowjacket ven
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RESULT 5
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hypothetical protein ZK384.2 - Caenorl
C;Species: Caenorhabditis elegans
C;Date: 15-Oct-1999 #sequence_revision
C;Accession: T27834
R;Ainscough, R.
submitted to the EMBL Data Library, No
A;Reference number: Z20427
A;Reference number: Z20427
A;Accession: T27834
A;Status: preliminary; translated from
A;Molecule type: DNA
A;Cross-references: UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:062508; UNIPROT:0
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A;Molecule type:
A;Residues: 1-241
A;Cross-reference
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C;Date:
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A; Gene:
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A; Map po
A; Intro
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;Introns: 60/3; 106/3; 180/2
;Superfamily: yellowjacket venom allergen antigen
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Similarity 29.0%; Pred. No. 1.1e-06;
47; Conservative 27; Mismatches 50;
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Similarity 24.7%; Pred. No. 1.6
6; Conservative 28; Mismatches
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PNCLCDFRFGPTAQREIVDFHNSLRSQLANGDYVV-
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   LEEVAAQQIAPCNDPLPINTSLAQN-
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RESULT 7
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hypothetical protein F49E11.4 - Cae
C; Species: Caenorhabditis elegans
C; Date: 15-Oct-1999 #sequence_revis:
C; Accession: T22439
R; Baynes; C.
submitted to the EMBL Data Library,
A; Reference number: Z19564
A; Accession: T22439
A; Status: preliminary; translated f:
A; Molecule type: DNA
A; Residues: 1-213 <WIL>
A; Cross-references: UNIPROT:Q93747;
A; Experimental source: clone F49E11
C; Genetics:
A; Gene: CESP:F49E11.4
A; Introns: 14/1; 60/3; 105/3; 175/2
C; Superfamily: yellowjacket venom a
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C; Species: Caenorhabditis el
C; Date: 15-Oct-1999 #sequenc
C; Accession: T22438
R; Baynes, C.
submitted to the EMBL Data I
A; Reference number: Z19564
A; Accession: T22438
A; Status: preliminary; trans
A; Residues: 1-207 <WIL>
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RESULT T19852 hypothe C; Speci C; Date:

tical protein C39E9.2 - Caenorha
es: Caenorhabditis elegans
15-Oct-1999 #sequence_revision

Caenorhabditis elegans

15-Oct-1999

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A;Status: p
A;Molecule
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C;Specie
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submitte
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A;Cross-references: UNIPROT:Q20603; UNIPARC:UPI0000
A;Experimental source: clone F49E11
C;Genetics:
A;Gene: CESP:F49E11.10
A;Map position: 4
A;Introns: 58/3; 104/3; 175/2
C;Superfamily: yellowjacket venom allergen antigen
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es: Caenorhabditis elegans
15-Oct-1999 #sequence_rev:
sion: T22432
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Local Similarity 23.0%;
es 56; Conservative 2
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ence number: Z19564
sion: T22432
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IGLC
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                                                                                                                                                    SWYWVSASLGFMKGTKLDQFANQ---
                                                         KSNLIGCGVKDCGRDSNGLNKVTVVCQYKPQGNFINQYIYVSG--ATCSGCPSGTSCETS
                                                                                        RNRKVGCAHKICPAQQN-----MVVSCVYGSPKLAPNEVIWQEGKACVCDARPDSFCCD--
                                                                                                                     YWYWTSGSLG-----DLNQYGSAASASWEKEFQDYGWKSNLMTIDLFNTGIGHATQMAWA
                                                                                                                                                                                                                                                                             VVVAVLLALFSYAEAGFCCPNSLSQSDSARQIFLDFHNDVRRNIALGNGLINWTVNADAV
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Pred. No. 0.0025;
25; Mismatches 89;
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submitted to the EMBL Data Librar, A; Reference number: Z19187
A; Recession: T19852
A; Status: preliminary; translated A; Molecule type: DNA
A; Residues: 1-208 <WIL>
A; Cross-references: UNIPROT:Q1854
A; Experimental source: clone C39E
C; Genetics:
A; Gene: CESP:C39E9.2
A; Introns: 14/1; 59/3; 176/2
A; Introns: 14/1; 59/3; 176/2
C; Superfamily: yellowjacket venom
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hypothetical protein F02E11.
C;Species: Caenorhabditis el
C;Date: 29-Oct-1999 #sequenc
C;Accession: T31959
R;Favello, A.; Scheet, P.
submitted to the EMBL Data L
A;Description: The sequence
A;Reference number: Z21104
A;Accession: T31959
A;Status: preliminary; trans
A;Molecule type: DNA
A;Residues: 1-207 <FAV>
A;Cross-references: UNIPROT:
A;Experimental source: strai
C;Genetics:
A;Gene: CESP:F02E11.5
A;Map position: 2
A;Introns: 59/3; 175/2
C;Superfamily: yellowjacket
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;Species: Caenorhabditis elegans
;Date: 29-Oct-1999 #sequence_rev;
;Accession: T31959
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15-Oct-1999 #sequence_revision 15-Oct-1999
sion: T19859; T22441
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ues: 1-211 <WIL>
-references: UNIPROT:Q18549;
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                                                                                                                                                                   VVVAVILALFSYAEAGFCCPNSLSQSDSARQIFLDFHNDVRRNIALGNGLINWTVNADAV
                                                                       ETVLQQVSWYWVSASLGFMKGTKLDQFANQWAEPLANIANYRNRKVGCAHKIC---
                                                                                                                     ILGPAQNMYKVDWDCNLEEVAAQQIAPCNDPLPINTSLAQNIA-----RW--LYFKDSEE
                                                                                                                                              LLVLVTLQIGAYAQF-
NYNRAVVVCQYKAQGNYLGQDIYKSG--TTCSACPTGTTCEAATGLC
                                             YGGAASVAWEQEFQQYGWTTNTFTQALANTGIGHATQMAWANTGLIGCGVKNCGPDPELN
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yellowjacket
                                                                                               --AGSNLLKMKWDTTL-ATAAQTFA---NTCPRGHSNAAGVGENLYWRWSSLPFSGMDI
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                          --MVVSCVYGSPKLAPNEVIWQEGKACVCDARPDSFCCD---NLC
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Z19187
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Pred. No. 0.0038;
5; Mismatches 107
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A; Title:
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    ; Accession:
                                                                               lar protein Tpx-1 - rat
les: Rattus norvegicus (Norway rat)
: 21-Aug-1998 #sequence_revision 21-Aug-1998
ssion: JE0204
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  ence number:
sion: JE0204
                T.; Sakashita, M.; Ohba, Y.; Nakanishi, Y. Biophys. Res. Commun. 248, 140-146, 1998 Molecular cloning of the rat tpx-1 responsible nce number: JE0204; MUID:98340864; PMID:9675100
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hypothetical protein C39E9.6 - Caenor C; Species: Caenorhabditis elegans C; Date: 15-Oct-1999 #sequence_revisio: C; Accession: T19849
R; Sims, M.
submitted to the EMBL Data Library, MA; Reference number: Z19187
A; Recession: T19849
A; Status: preliminary; translated from A; Molecule type: DNA
A; Residues: 1-210 <WIL>
A; Cross-references: UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q18540; UNIPROT:Q1
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A; Gene: CESP: C39E9.6
A; Map position: 4
A; Introns: 58/3; 178/2
C; Superfamily: yellowja
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C; Comment: This pro
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A; Map position: 17
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C; Superfamily: yellowjacket vo
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C;Species: Caenorhabditis elegans
C;Date: 15-Oct-1999 #sequence_revision 15-Oct-1999 #text_change
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R;Baynes, C.
submitted to the EMBL Data Library, March 1996
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OM nucleic - nucleic search, using sw model Run on: March 4, 2006, 07:04:30; Search time; 4211 Seconds [without alignments] 10299:585 Million cell updates/sec Title: 19-10-527-771-9 10299:585 Million cell updates/sec Title: 19-10-527-771-9 Sequence: 1 98-8actgctatgtcggcggtcatgcasasasasasa 763 Secoring table: IDENTITY NUC Gapop 10-0, Gapext 1.0 Searched: 5883141 seqs, 28421725653 residues Total number of hits satisfying chosen parameters: 11-95-10: 11-95-1
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/gene="adspz" /codon_start=1 /product="ancyclostoma-secreted protein-like protein" /protein_id="CAD56659.1" /db_xref="GI:25005300" /db_xref="GOA:Q8IFT6" /db_xref="UniProt/Trembl:Q8IFT6" /translation="MSAAVVVAVLLALFSYAEAGFCCPNSLSQSDSARQIFLDFHNDV	11706	1763	<pre>/organism="Ostertagia ostertagi" /mol_type="mRNA" /db_xref="taxon:6317" /dev stage="adult"</pre>	1763	University, Salisburylaan 133 Merelbeke, 9820, BELGIUM Location/Qualifiers	07-NOV-2002) Geldhof P.B., Parasito	Direct Submission	eldhof.P.B.	2 (bases 1 to 763)	MOI. DIOCHEM. Farabicoi. 120 (1), iii-iii (2003)	ens in a host protective frac	ociated secreted	ove, J., Vercruysse, J. and Claerebout, E.	Geldhof, P., Vercauteren, I., Gevaert, K., Staes, A., Knox, D.P.,	Tricnostrongyloidea; Haemonchidae; Ostertagiinae; Ostertagia. 1	Eukaryota; Metazoa; Nematoda; Chromadorea; Rhabditida; Strongylida;	ostertagi	agia ostertagi	٠.	523.1	procein (aaspz gene). AJ515523	agia ostertagi mRNA for ancylostoma-secreted prote	OOS515523 763 bp mRNA linear INV 18-APR-2003		

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Maere, V.
Ostertagia vaccine
Patent: WO 2004024769-A 9
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GCGAGATGCTGCGAGTGTTCGCCACCAGTGTTGCGCGTCGCCATGAAGCGAAAAGAAATT
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Patent: WO 20
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Trichostrongyloidea; Haemonchidae; Osterta
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mol_type="unassigned
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Submitted (07-FEB-2002) Vercauteren I.J.R., Department
Parasitology, Ghent University, Faculty of Veterinary I
Salisburylaan, 133, B-9820 MERELBEKE, BELGIUM
On Feb 22, 2002 this sequence version replaced gi:13530
Location/Qualifiers
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Direct Submission
Submitted (13-AUG-2001) Vercauteren I.J.R., Department
Parasitology, Ghent University, Faculty of Veterinary |
Salisburylaan, 133, B-9820 MERELBEKE, BELGIUM
revised by [4]
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Vercruysse, J.
Vercruysse, J.
Identification of excretory-secretory products of
Ostertagia ostertagi by immunoscreening of cDNA 15
Mol. Biochem. Parasitol. 126 (2), 201-208 (2003)
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Vercauteren, I.J.R.
Direct Submission
Submitted (29-MAR-2001) Vercauteren I.J.R., Department of
Parasitology, Ghent University, Faculty of Veterinary Medicine,
Salisburylaan, 133, B-9820 MERELBEKE, BELGIUM
Revised by [3]
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/db_xref="GI:18860824"
/db_xref="GOA:Q8T8F4"
/db_xref="UniProt/TrEMBL:Q8T8F4"
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Chang, H.S., Chen, W., Cooper, B., Glazebrook, J., Goff, S.A., Hou
Katagiri, F., Quan, S., Tao, Y., Whitham, S., Xie, Z., Zhu, T. and
Plant genes involved in defense against pathogens
Patent: WO 03000898-A 5263 03-JAN-2003;
Syngenta Participations AG (CH)
Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /codon_start=1
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/protein_id="BAD16466.1"
/protein_id="BAD16466.1"
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AVAATKGYNVQIVVDENEGEESIFRRFREVMRAGVLQEIKRRRYESKKDERKRKAR
EAGRRNRRRRMMEEPFFPEDDADSEAARRADDEKDNWEIDGIL"
complement(join(9908. .13128)
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complement(join(9908. .10162,11120. .11325,12777. .12893,
13018. .13128))
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/contains full-length cDNA(s): AK111066
similar to Arabidopsis thaliana chromosome 3, At3g01435"
/codon_start=1
/product="unknown protein"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /note="copia-type polyprotein-like"
complement (7653. .8923)
/gene="OSJNBb0031B09.3"
complement (join (7653. .7929,8479. .89
/gene="OSJNBb0031B09.3"
/note="supported by full-length cDNA(
complement (join (7821. .7929,8479. .88
/gene="OSJNBb0031B09.3"
/note="contains EST(8): AU161887(S619)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /gene="OSJNBb0031B09.5"

join(13885. .14326,14395. .15378)

/gene="OSJNBb0031B09.5"

/note="supported by full-length cDN
join(13898. .14326,14395. .15099)

/gene="OSJNBb0031B09.5"

/note="contains full-length cDNA(8)
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RLRSFRGDDDAAAAAAATLSEALYIVSMGTNDFLENYYAVARGHAAEYSTAAAYGDY
                                                                                                                                                                                                                              LLGVAEAFVRELHALGARKVDLNGLPPMGCLPLBRATGGACTEEYNAVAGRFNAGLQD
MIARLNGELGGGARIVYGDVYGAVAAVLADPAAYGVENVKAGCCGVTGVFEMGYMCGA
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/db_xref="GI:46390954"
/translation="MLPQSQSSSLQRLNHVEQMIVRAVNLAGTVMEELGNATGPRTEG
VAGHCREFMLAMKEIQTTLREEIKSACEYRPFEKCDYSARIANBICCKKVEYVLEKLD
                                                                                                                                                                                                                                                                                                                                                                                                                                        precursor"
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RRGGARGARGRRDGRAATATADGDGGSGGARREAWSTGAATRRPRGDGDGMAARRRRW
RQAARHGARAAGARGSVMATAGSGGTGLGAVSARQWGRVRLGNLVEHLAADB"
join(2424...2428,3888...4800,6598...6753,6862...7095)
                                                                                                                                                                                                            15811.
   /codon_sta
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13885. .15378
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                                                                  note="supported by full-
oin(15882. .16277,17884.
gene="OSJNBb0031B09.6"
                                                                            /gene="OSJNBb0031B09.2"
|oin(2424. .2428,3888. .4800,
|gene="OSJNBb0031B09.2"
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                                                  note="contains
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db_xref="GI:46390955"
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NA(Β): AK120425"
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                                                  CDNA(8):
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VERSION KEYWORDS SOURCE

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AC131303 AC131303.8

GI:51854578

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                                                                                                                                                 AAGGCACGAAACTTGACCAATTTGCTAACCAGTGGGCTGAACCTCTAGCAAACATTGCAA
                                                                                ACTATAGAAACCGAAAGGTTGGATGTGCCCCATAAG
                                                                                                                  TTGGAGCAGAACTAATCGATTTTACCTÁCAATTGGCTTGTAAATTTCACAAGCATTGCAA
                                                                                                                                                                                     AAGAGACAGTTCTGCAACAAGTATCGTGGTATTGGGTGAGCGCATCGCTGGGATTTATGA
                                                  <u>AATGTGAAATTCAACATTCTCGGCTTTCAAATATG</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     complement (20777. .23474)
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complement (join(20777. .20936,21084. .21920,23153.
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contains full-length cDNA(s): AK108100"
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GYFPLLLQILRVTACSPARRSKASHVPDVAPVVCNSGDNDGDGYNNRSWALVMKLLSV
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VDTGNNNQIGTPLRSDFPPYGRDMPGGARATGRFGNGRLAPDFMSESLGLPPLVPAYL
DPAYGIADFARGVCFASAGTGLDNATAGVLSVIPLWKEVEYYREYQRRLRAHAGAAA
RDVVRGALHVVSIGTNDFLENYYMLATGRFARYSVGEYEDYLVAAARAFLAAIHRLGA
RRVTFAGLSPMGCLPLERTAGALLGGGGGGCVEEYNRVAREYNGKVEAMVRSLRAELP
RLKVAFIPVYDNMLDLITHPEKYGLENVEEGCCATGRFEMGFMCNDESPLTCDDASKY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               complement (25261. .26644)
/gene="OSJNBb0031B09.8"
complement (join (<25261. .25446,26001.
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/note="start and end point are not id.
complement (join (25261. .25446,26001.
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29116. .29268
                                                                                                                                                                                                                                                                                                                                                                             29116.
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                                                                                                                                                                                                                                                                                                                            ="OSJNBb0031B09.9"
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                                                                                                                                                                                                                                                                                                                                                                               29268
                                                                                                                                                                                                                                                      0;
                                                                                                                                                                                                                                                                      Score 41.4;
Pred. No. 0.
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NA(s): AK108100"
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REFERENCE
AUTHORS
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JOURNAL
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2 (Nabes 1 to 187110)
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8 Birren, B., Nubbaum, C., Lander, E., Ali, A., Allen, N., Anderson, S., Birren, B., Nubbaum, C., Lander, E., Ali, A., Allen, N., Bageler, B., Callymore, A., Choepel, Y., Collymore, A., Barne, N., Bageler, J. Chang, J., Chazaro, H., Choepel, Y., Collymore, A., Cook, A., Cooke, P., Dekrellano, K., Dewar, K., Galagan, J., Chazaro, P., Pitcaferald, M., Gege, D., Galagan, J., Camarata, J., Chang, P., Pitcaferald, M., Gege, D., Galagan, J., Cook, A., Cooke, P., Dekrellano, K., Dewar, K., Galagan, J., Charlen, C., Macchand, P., Major, J., Matthews, C., Kamata, S., Gord, S., Graham, L., Grand-Pierre, N., Hagos, B., Karatas, A., Kalls, C., Landers, T., Levine, R., Lindblad-Toh, K., Liu, G., MacLean, C., Macchand, P., Major, J., Matthews, C., McCarthy, M., Peldrim, J., Meneus, L., Mihova, T., Mienga, V., Marphy, T., Naylor, J., Nguyan, C., Netta, R., Rise, C., Rogov, P., Romar, J., Roy, A., Schauer, S., Schupback, R., Seaman, S., Severy, P., Phunkhang, P., Pierre, N., Raymond, C., Retta, R., Rise, C., Rogov, P., Romar, J., Roy, A., Schauer, S., Schupback, R., Seaman, S., Severy, P., Phunkhang, P., Pierre, N., Haynond, C., Retta, R., Rise, C., Rogov, P., Romar, J., Roy, A., Schauer, S., Schupback, R., Seaman, S., Severy, P., Smith, C., Spencer, B., Stange-Thomann, N., Scojanovic, N., Talamas, J., Traffer, S., Schupback, R., Seaman, S., Severy, P., Smith, C., Spencer, B., Stange-Thomann, N., Scojanovic, N., Talamas, J., Tander, B., Stange-Thomann, N., Scojanovic, N., Talamas, J., Stange-Thomann, N., Scolanovic, N., Talamas, J., Stange-Thomann, N., Scolanovic, N., Talamas, J., Stange-Thomann, N., Stange-Thomann, N
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4 (bases 1 to 187110)

S Birren, B., Nusbaum, C., Lander, E., Abouellell, A., ...

Anderson, M., Anderson, S., Arachchi, H.M., Barna, N., Bastien, V., Anderson, M., Anderson, S., Arachchi, H.M., Barna, N., Bastien, V., Anderson, M., Anderson, S., Arachchi, H.M., Barna, N., Chang, J., Chong, T., Boguslavkiy, L., Boukhgalter, B., Commarata, J., Chang, J., Choepel, Y., Collymore, A., Cook, A., Cooke, P., Corum, B., DeArellano, K., Diaz, J.S., Dodge, S., Dooley, K., Dorris, L., DeArellano, K., Diaz, J.S., Dodge, S., FitzGerald, M., Gage, D., Erickson, J., Gardyna, S., Graham, L., Grand-Pierre, N., Hafez, N., Hagopian, D., Hagos, B., Hall, J., Horton, L., Hulme, W., Iliev, I., Johnson, R., Jones, C., Kamat, A., Karatas, A., Kells, C., Landers, T., MacLean, C., Macdonald, P., Major, J., Manning, J., Matthews, C., McCarthy, M., Meldrim, J., Meneus, L., Mihova, T., Micol, R., Norbu, C., McCarthy, M., Maylor, J., Nguyen, C., Nguyen, T., Nicol, R., Norbu, C., O'Neil, D., Oliver, J., Peterson, K., Dodge, C., Macthews, 
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Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia;
Sciurognathi; Muroidea; Muridae; Murinae; Mus.
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Birren, B., Nusbaum, C. and Lander, E.
Mus musculus chromosome 1, clone RP24-119019
Raymond, C., Schupback, R.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Contact: sequence submissions@broad.mit.edu
----- Project Information
Center project name: L27079
Center clone name: 119_0_19
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Web site: http://www-seq.wi.mit.edu
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4395. .4519

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2348. .2378
/rpt_family="AT_rich
2844. .3688
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3680. .4391
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5609. .5923
/rpt_family="Lx8"
7795. .7881
/rpt_family="(TA)n"
7889. .7938
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8421. .8504
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385.
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1932. .2018
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1016. .1251
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688. .713
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10446. .10649
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8324._.8420
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mol_type="genomic DNA"
db_xref="taxon:10090"
chromosome="1"
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clone_lib="RPCI-24
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complement(26204. .26233)
/rpt_family="B3A"
26234. .26267
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21578.
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lement(18812. .21567)
family="LIVL2"
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7. .18265
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7. .17657
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ement(21745. .21880)
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y="RLTR11A"
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                                                                                                 Sequencing of Mus muscult Unpublished (2001)
3 (bases 1 to 198900)
McPherson, J.D. and Waters Direct Submission
Submitted (30-MAY-2002) (Parkway, St. Louis, MO 634 (bases 1 to 198900)
McPherson, J.D. and Waters Direct Submission
Submitted (11-FEB-2003) (Submitted (11-FEB-2003) (51)
                                                                                                                                                                                                                                                                                                                                                                                                                            Sciurognathi; Muroidea; Muridae;
1 (bases 1 to 198900)
Dauphin,S., Meyer,R., Kohlberg,S
The sequence of Mus musculus BAC
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AC123556
AC123556.4 GI:29244829
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                                                                                                                                                                                                                                                                                                                                                                                                                   Unpublished
                              Center: Washington University Genome Center code: WUGSC
Web site: http://genome.wustl.edu
Contact: submissions@watson.wustl.edu
----- Summary Statistics
Center project name: M_BA0176M09
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usculus BAC clone RP23-176M9
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clone. It may be shorter because we only sequence overlapping clone sections once, or longer because we provide a small overlap between neighboring data submissions.

Inis sequence was finished as follows unless otherwise noted: all regions were double stranded, sequenced with an alternate chemistry, or covered by high quality data (i.e., phred quality >: 30); an attempt was made to resolve all sequencing problems, such as compressions and repeats; all regions were covered by sequence from more than one subclone; and the assembly was confirmed by restriction digest.

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MAPPING INFORMATION:
Mapping information for this clone was provided by Dr.
Department of Genetics, Washington University, St. Lou
additional information about the map position of this http://genome.wustl.edu Dr. We Louis sequence, ₹ Ö Warren, Por вее

INFORMATION:

The RPCI-23 BAC Library has been constructed by Kazutoyo Oseg and Minako Tateno in the laboratory of Pieter de Jong (http://www.chori.org) from female C57BL/6J mouse kidney and/brain genomic DNA. The clone and detailed information can be obtained from Research Genetics, Inc. (http://www.resgen.com) Pieter de Jong and coworkers at http://www.chori.org kidney and/or Osegawa or

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/rpt_family="ERV1"
5620. .5822
/rpt_family="L1"
6086. .6363
/rpt_family="ERVK"
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/rpt_family="ERVK"
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7816. .7956
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/rpt_family="B2"
13518. .13601
/rpt_family="B4"
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/rpt_family="L1"
15729. .16048
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chromosome="1"
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clone_lib="RPCI-23"
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Cambridge, MA 02141, USA
http://www-seq.wi.mit.edu
This sequence was finished as follows unless otherwise not regions were either double-stranded or sequenced with an acchemistry or covered by high quality data (i.e., phred quality an attempt was made to resolve all sequencing problem as compressions and repeats; all regions were covered by one subclone; and the assembly was confirmed by restriction except on the rare occasion of the clone being a YAC.

Location/Qualifiers
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Leongamornlert, D.

Direct Submission

Submitted (13-MAY-2005)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Web site: http://www.sanger.ac.uk
Contact: vega@sanger.ac.uk
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Mammalia; Eutheria;
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/db_xref="taxon:9606"
/chromosome="6"
/clone="RP11-173D14"
/clone_lib="RPCI-11.1"
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A sequence from clone RP11-173D14 on chromosome 6 Contains
a novel gene, a gene for a novel protein (likely ortholog
putative IKK regulator SIMPL), the 3'end of a gene for a
repeat-containing protein and a CpG island, complete
99396. .130527
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Center: Washington University Genome Center code: WUGSC
Web site: http://genome.wustl.edu
Contact: sapiens@watson.wustl.edu
Contact: Summary Statistics
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NOTICE: This sequence may not represent the entire insert of this clone. It may be shorter because we only sequence overlapping clone sections once, or longer because we provide a small overlap between neighboring data submissions.

This sequence was finished as follows unless otherwise noted: all regions were double stranded, sequenced with an alternate chemistry, or covered by high quality data (i.e., phred quality >= 30); an attempt was made to resolve all sequencing problems, such as compressions and repeats; all regions were covered by sequence from more than one subclone; and the assembly was confirmed by restriction digest.

MAPPING INFORMATION:

The sequence of this clone was established as part of a mapping and sequencing collaboration between the NHGRI Chromosome 7 Mapping Project (Bric D. Green, Director), John D. McPherson in the Department of Genetics (Washington University), and the Washington University Genome Sequencing Center. For additional information about the map position of this sequence, see http://www.nhgri.nih.gov/DIR/GTB/CHR7, send mailto:egreen@nhgri.nih.gov, or see http://genome.wustl.edu

SOURCE INFORMATION:
The RPCI-11 human BAC library was made from the blood of one donor, as described by Osoegawa, K., Woon, P.Y., Zhao, B., Freng Tateno, M., Catanese, J.J. and de Jong, P.J. (1998) An improved approach for construction of bacterial artificial chromosome libraries. Genomics 51:1-8. The clone may be obtained either Research Genetics, Inc. (http://www.resgen.com) or Pieter de and coworkers at http://www.chori.org f one male Frengen, E., from Jong

NEIGHBORING The clone se BORING SEQUENCE II lone sequenced to the INFORMATION: to the left i he right is R is RP11-468B6, 200 bp RP11-561N12. Actual p overlap start of the this

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27297..27572
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Marie, Merzker M.Lee., Abramzon, S., Adams, C., Alder, J., Sharry, D., Marie, Merzker, M.Lee., Abyaer, A., Angulano, D., Alder, J., Alabrooke, S., Anin, A., Angulano, D., Anyalabechi, V., Acyagi, A., Ayodeji, M., Baca, B., Baden, H., Mayalabechi, V., Acyagi, A., Ayodeji, M., Baca, B., Baden, H., Barley, B., Baden, H., Cardena, V., Henderson, N., Henderson, N., Henderson, N., Henderson, V., Hennedea, J., Hernandez, V., Havlak, P., Hawes, A., Henderson, N., Kovar, C., Kowis, C., Lake, J., S., Kally, S., Khan, Z., King, L., Kovar, C., Kowis, C., Lake, J., J., Liu, Y., London, P., Longocre, S., Lopez, J., Liu, J., Liu, Y., London, P., Longocre, S., Lopez, J., Liu, J., Liu, Y., London, P., Longocre, S., Lopez, J., Liu, J., Liu, Y., London, P., Longocre, S., Lopez, J., Hunderson, J., Henderson, N., Martin, R., M
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1 (bases 1 to 225475)
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On Nov 19, 2002 this sequence version replaced gi:22856233.
The sequence in this assembly is a combination of BAC based reads and whole genome shotgun sequencing reads assembled using Atlas (http://www.hgsc.bcm.tmc.edu/projects/rat/). Each contig described in the feature table below represents a scaffold in the Atlas assembly (a 'contig-scaffold'). Within each contig-scaffold, individual sequence contigs are ordered and oriented, and separated by sized gaps filled with Ns to the estimated size. The sequence contigs within a contig-scaffold that consist entirely of whole genome shotgun sequence reads. Both end sequences and whole genome shotgun sequence only contigs will be indicated in the feature
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113; Conservat
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Direct Submission
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* of the gaps between them are bas

* provided by the submittor.

* This sequence will be replaced

* by the finished sequence as soon

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1 225475: contig of 225

Location/Qualifiers

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Center project Information
Center project name: GWVW
Center clone name: CH230-25B23
Center clone name: CH230-25B23
Rssembly program: Phrap; version 0.990329
Consensus quality: 206303 bases at least Q40
Consensus quality: 208669 bases at least Q30
Consensus quality: 210148 bases at least Q20
Estimated insert size: 212963; sum-of-contigs estimation
Quality coverage: 6x in Q20 bases; sum-of-contigs estimation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Center code: BCM
Web site: http://www.hgsc.bcm.tmc.edu/
Contact: hgsc-help@bcm.tmc.edu
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                                                                                                                                                                                                                                                                                                                                                                                           http://www.hgsc.bcm.tmc.edu/docs/Genbank_draft_data.html).
This is a 'working draft' sequence. It currently
sof 1 contigs. Gaps between the contigs
resented as runs of N. The order of the pieces
eved to be correct as given however.
                                                                                                          /note="wg8_co
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                                                                                                                                                                organism="Rattus norvegicus"

mol_type="genomic DNA"

/db_xref="taxon:10116"

/clone="CH230-25B23"
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d. No. 3.6; Mismatches
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                                                                                                                                                                                                                      Muzny, D. Marie, Metzker, M. Lee, Abramzon, S., Adams, C., Alder, J., Allen, C., Allen, H., Alsbrooks, S., Amin, A., Anguiano, D., Allen, H., Alsbrooks, S., Amin, A., Anguiano, D., Allen, H., Alsbrooks, S., Amin, A., Anguiano, D., Allen, H., Aoyagi, A., Ayodeji, M., Baca, E., Baden, H., Baldwin, D., Bandaranaike, D., Barber, M., Barnstead, M., Benahmed, P., Baldwin, D., Bandaranaike, D., Barber, M., Barnstead, M., Benahmed, P., Baldwin, D., Bandaranaike, D., Barber, M., Barnstead, M., Benahmed, P., Baldwin, D., Bandaranaike, D., Barber, M., Barnstead, M., Benahmed, P., Burch, P., Burreil, K., Calderon, E., Chu, J., Chen, S., Durland, C., Chen, Y., Chen, Z., Chu, J., Checko, J., Charter, K., Cavazo, I., Chen, R., Chen, Z., Chu, J., Checko, J., Charter, R., Cox, C., Coyle, M., Cree, A., D., Souza, L., Carden, R., Chen, Y., Chen, Z., Chu, J., Checko, J., Chen, S., Durland, R., Cree, A., D., Souza, L., Cheland, C., Dederich, D., Davila, R., Davila, R., Chen, Z., Chu, J., Chen, G., Chen, Y., Chen, Z., Chu, J., Chen, S., Durland, R., Durland, C., Dederich, D., Davila, R., Davila, R., Carder, M., Foster, P., P., Farls, C., Chen, S., Durland, C., Dederich, D., Davila, R., Davila, R., Carde, A., Garrer, T., Garzá, M., Foster, P., Farser, C.M., Gabisi, A., Ganta, R., Garcia, A., Garrer, T., Garzá, M., Gebregeorgis, E., Geer, K., Gill, R., Grady, M., Guerra, W., Guevara, W., Harvey, Y., Hawes, A., Henderson, N., Hernandez, J., Hernandez, J., Harvey, Y., Havlak, P., Hawes, A., Henderson, N., Hernandez, J., Jackson, A., Hollins, B., Howells, S., Hullyk, S., Hume, J., Idebird, D., Jackson, A., Jackson, A., Jackson, A., Hollins, B., Hoyels, S., Haldun, S.L., Hodgson, A., Hogues, M., Harvey, Y., Havlak, P., Johnson, B., Johnson, R., Jolivet, A., Kowis, C., Liu, J., Liu, W., Liu, Y., London, P., Longacre, S., Lopez, J., Liu, J., Liu, W., Liu, Y., London, P., Longacre, S., Lopez, J., Liu, J., Liu, W., Liu, Y., London, P., Longacre, S., Lopez, J., Liu, J., Liu, W., Liu, Y., London, P., Longacre, S., Lopez, J., Liu, J., Liu, W., L
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                                            Pasternak, S., Paul, H., Perez, A., Perez, L., Pfannkoch, C
Plopper, F., Poindexter, A., Popovic, D., Primus, E., Pu, L
Puazo, M., Quiroz, J., Rachlin, E., Reeves, K., Regier, M.A
Reilly, B., Reilly, M., Ren, Y., Reuter, M., Richards, S., l
Rives, C., Rodkey, T., Rojas, A., Rose, M., Rose, R., Ruiz, Sanders, W., Savery, G., Scherer, S., Scott, G., Shatsman, Shetty, J., Shvartsbeyn, A., Sisson, I., Sitter, C.D., Sman, Sneed, A., Sodergren, E., Song, X.-Z., Sorelle, R., Sosa, J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        HTG; HTGS_PHASE1;
Rattus norvegicus
Rattus norvegicus
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Sodergren, E., Sulton, A., Sutton, A.,
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thi; Muroidea;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Eutheria; Euarchontoglires; Glires; Rodentia;
hi; Muroidea; Muridae; Murinae; Rattus.
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Svatek, A., law-
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, WORKING DRAFT SEQUENCE, 3
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us,E., Pu,L.-L.,

Regier,M.A., Reigh,R

Regies, Riggs,F.,
      R., Sosa,J.,
Tabor,P., Trejos,Z., (
                                                                                              Smaje, D.
          Taylor, C., Usmani, K.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Foster, P.,
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                                                                                                                           Shen, H.
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On Nov 6, 2002 this sequence version replaced gi:23908010.
The sequence in this assembly is a combination of BAC based reads and whole genome shotgun sequencing reads assembled using Atlas (http://www.hgsc.bcm.tmc.edu/projects/rat/). Each contig described in the feature table below represents a scaffold in the Atlas assembly (a 'contig-scaffold'). Within each contig-scaffold, individual sequence contigs are ordered and oriented, and separated by sized gaps filled with Ns to the estimated size. The sequence contigs within a contig-scaffold that consist entirely of whole genome shotgun sequence reads. Both end sequences and whole genome
                   _feature
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Direct Submission
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Direct Submission
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Center clone name: CH230-415F5
Center clone name: CH230-415F5
Assembly program: Phrap; version 0.990329
Consensus quality: 172958 bases at least Q40
Consensus quality: 175322 bases at least Q30
Consensus quality: 176115 bases at least Q20
Estimated insert size: 174254; sum-of-contigs estimation
Estimated insert size: 174254; sum-of-contigs estimation
Consensus quality: 176115 bases; sum-of-contigs estimation
Estimated insert size: 174254; sum-of-contigs estimation
Consensus quality: 176115 bases; sum-of-contigs estimation
Estimated insert size: 174254; sum-of-contigs estimation
Consensus quality: 176115 bases; sum-of-contigs estimation
                                                                                                                                                                                                                                                                                                                  NOTE: Estimated insert size may differ from sequence 16 (see http://www.hgsc.bcm.tmc.edu/docs/Genbank_draft_NOTE: This is a 'working draft' sequence. It currently consists of 3 contigs. The true order of the pieces is not known and their order in this sequence record is arbitrary. Gaps between the contigs are represented as runs of N, but the exact sizes of the gaps are unknown. This record will be updated with the finished sequence as soon as it is available and the accession number will
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236396
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                                                /mol_type="genomic DN
/db_xref="taxon:10116
/clone="CH230-415F5"
                                                                                                                                                             Locat
                                               clone="CH23
                                                                                                               organism="Rattus
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237724: contig of 237824: gap of unka
239001: contig of 239001: contig of ation/Qualifiers
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VERCRUYSSE J.
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Pred. No. 7.6e-76;
3; Mismatches 244;
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one gene involved in plant resistance or response for conferring resistance or tolerance to a plant viral infection by determining or detecting plant
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Matches
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Similarity 9.3%; Pred.
64; Conservative 263; M
KMWRRYATRMMWMWYRYSMKWYTWCTMWGYWWYWWRTYMKMRY
                                                                    HISGSRMSRWKSYTCYWRKWGSMKSTCTWMYYMSKYTYAKYGSYWRYRYRAWCMYMWRWY
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The invention relates to an isolated nucleic acid comprising at least 10 contiguous nucleotides of any of the 233 polynucleotide sequences given in the specification, or its complement. The nucleic acids encode cancerassociated proteins. Also included are an expression vector comprising the above recombinant nucleic acid cited above, a host cell comprising the above recombinant nucleic acid or expression vector, a microarray for detecting recombinant nucleic acid or expression vector, a microarray for detecting comprising at least 10 contiguous nucleotides of any of the above comprising at least one probe comprising and the above comprising the above antibody and a pharmaceutical composition comprising the above colls in an individual, a method for inhibiting growth of cancer cells in an individual, an electronic library comprising the above concer cells in an individual, an electronic library comprising the above cells in an individual, an electronic library comprising the above cells in an individual, an electronic library comprising the appreciating at methods for detecting cancer of method for tracting and a method for inhibiting the expression of comprisions and a method for lamble, and cell sample, a comprision of the probability of compris
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                                 pecification, but was obtained in electron
ftp.wipo.int/pub/published_pct_sequences
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ukemia, or in
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Isolated nucleic acid molecules having polymorphisms in e.g. cytochrome p450 and cathepsin S useful as genetic for locating, identifying and characterizing the genes
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This invention relates to the sequence of an isolated mucleic acid molecule comprising at least one base variation from that of a known busine p450 A1 (CYP4501A1), cytochrome P450 A2 (CYP4501A2), cytochrome P450 A2 (CYP450
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\$ 멍 8 S ঠ S 멍 Query Ma Best Loc Matches Local Match 642 522 CCGCTCAGCAAAACATGGTAGTATCCTGCGTGTATGGAAGCCCCAAACTTGCACCGAACG 581 111; Similarity GCGACAACCTGTGTGACACGCGAGATGCTGCGAGTGTTCGCCACCAGTGTTGCGCGTCGC 701 TAAAAAATTAGCCAGACGCAGTGCGTGTGCCTGCAATCCCCAGCTACTCGTGAGACTGAG 794 GCAGGAGAATTGCTTAAATCCGGGAGGCGGAGGTTGCAATGAGCCAAGATTGCGCCACTG Conservative 46.8%; Score 35.4; DI Pred. No. 4.7; O; Mismatches DB 6; Length 1537; 126; Indels 0; Gaps 0

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contiguous nucleotides of any of the 233 polynucleotide sequences given in the specification, or its complement. The nucleic acids encode cancerassociated proteins Also included are an expression vector comprising the isolated nucleic acid cited above, a host cell comprising the above recombinant nucleic acid cited above, a host cell comprising the above a cancer-associated (CA) nucleic acid comprising at least one probe comprising at least 10 contiguous nucleotides of any of the above comprising at least 10 contiguous nucleotides of any of the 95 polynucleotide sequences, an isolated polypeptide (encoded within an open reading frame of a CA sequence selected from any of the 95 polynucleotide sequences as mentioned in the specification, or its polynucleotide sequences as mentioned in the specification, or its polynucleotide shove polypeptide, a hybridoma that produces the above antibody and a pharmaceutical excipient, a kit for detecting the above monoclonal antibody, a pharmaceutical composition comprising the above antibody and a pharmaceutical excipient, a kit for detecting cancer cells (comprising the antibody cited above, methods for diagnosing cancer individual, a method for inhibiting growth of cancer cells in an individual, an electronic library comprising the above polynucleotide or polypeptide (or their fragments), methods of screening for associated with expression of a polypeptide in a test cell sample, a method for treating cancers and a method for inhibiting the expression of associated with expression of a polypeptide in test cell sample, a method for treating cancers and a method for inhibiting the expression of associated with expression of a polypeptide in a test cell sample, a method for treating cancers and a method for inhibiting the expression of leukaemia. These may also be used in screening for agents that modulate cancer. The present sequence is a mouse CAP genomic sequence. Note: The sequence data for this patent did not form part of the printed cancer for the printed sequence is a soc
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                                                  are associated with carcinomas. The sequences are useful for: (i) for screening drug candidates; (ii) for screening of bioactive agent capable of binding to Carcinoma Associated Protein (CAP); (iii) for screening of bioactive agent capable of modulating the activity of CAP; (iv) for carcinoma; (vi) for inhibiting the activity of CAP; (iv) for carcinoma; (vii) for neutralizing the activity of CAP; (ix) as a biochip; (c) for diagnosing carcinoma or a propensity to carcinoma; and (xi) for carcinoma or a propensity to carcinoma; and (xi) for carcinoma are useful as DNA vaccines and the CAP are useful as markers of carcinoma including lymphoma. The present sequence is one such CA coding customates and the carcinoma for the patent is an equivalent to basic patent customates.
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degree of hybridisation to both classes is determined from the label on the amplicon. From the ratio of labels hybridised to the two classes of oligomers, the degree of methylation is calculated. The method is used:

(i) for diagnosis and/or prognosis of side effects of therapeutic drugs and of a wide range of diseases, e.g. cancer, disorders of the central nervous, cardiovascular, gastrointestinal and respiratory systems etc., particularly by detecting mutations or single nucleotide polymorphisms (SNP's); and (ii) for differentiation of cell or tissue types and for investigating cell differentiation. The method allows the methylation status of many C residues to be determined simultaneously. ABQ13410-ABQ54121 represent genomic DNA sequences used to illustrate the method disclosure of the invention
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                                                                                                                                                                                                                                                                                                                                           This invention describes a novel nethylation of a particular cytogenomic sample of DNA. The sample cytosine (C) but not methylated on the target C is the amplicon is hybridised to two of oligonucleotides and/or peption of oligonucleotides and/or peption of hybridisation to both the contains the target C is a second to two of oligonucleotides and/or peptions of the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the contains the co
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(i) for diagnosis and/or prognosis of side effects of therapeutic drugs and of a wide range of diseases, e.g. cancer, disorders of the central nervous, cardiovascular, gastrointestinal and respiratory systems etc., particularly by detecting mutations or single nucleotide polymorphisms (SNP's); and (ii) for differentiation of cell or tissue types and for investigation of cell or tissue types and for
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RESULT 15
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Sequence

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B \$ B \$ 밁 Ş Query Match 4.5%; Best Local Similarity 51.3%; Matches 79; Conservative The invention relates to an isolated nucleic acid detection reagent capable of detecting 1000 or more genes from Drosophila. The invention useful in developmental biology and in elucidating cell signalling and cell-cell interactions in higher eukaryotes for the development of insecticides, therapeutics and pharmaceutical drugs. The invention discloses genomic DNA sequences (ABL16176-ABL30511), expressed DNA sequences (ABL01840-ABL16175) and the encoded proteins (ABB57737-ABB72072). The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences Sequence 1046 182 122 986 926 242 7160 BP; ACAGTAAATGCAGACGCGGTCATTCTTGGTCCAGCTCAGAACATGTACAAAGTGGACTGG CACGTATACCTCACTCAAGGAGATCGTACTAACCTTTTCGGAGTAGTTTGCATATGCAAG GATTGCAACTTGGAAGAAGTAGCAGCACAACAGA 275 CTCGATTTTCACAATGATGTTCGTCGAAAATATAGCACTTGGAAAATGGTTTGATAAACTGG 2075 A; 1468 C; Score 34; DB 4; Pred. No. 28; 0; Mismatches 1473 G; 2144 75; Ŧ; Length 7160; 0 Indels ų; 0 .Other; 0; Gaps 987 181 241). 19 0;

Search completed: March 4, 2006, 07:12:51 Job time : 594 secs

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APPLICATION NUMBER: BP 91 114 300.6
FILING DATE: 26-AUG-1991
ATTORNEY/AGENT INFORMATION:
NAME: BENT, Stephen A.
REGISTRATION NUMBER: 29,768
REFERENCE/DOCKET NUMBER: 30472/114 IM
TELECOMMUNICATION INFORMATION:
TELEPHONE: (703)836-9300
TELEFAX: (703)83-4109
TELEX: 899149
INFORMATION FOR SEQ ID NO: 14:
SEQUENCE CHARACTERISTICS:
LENGTH: 7218 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
IMMEDIATE SOURCE:
CLONE: pTZgpt-F1s

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	sequence seq
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ER OF SEQ ID NOS: 207012
WARE: FastSEQ for Windows 1
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Pred. No. 1.2
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                                                                         APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES AS:
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF
FILE REFERENCE: CL001307
CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
NUMBER OF SEQ ID NOS: 207012
SOFTWARE: FASTSEQ for Windows Version 4.0
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CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
NUMBER OF SEQ ID NOS: 207012
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 14790
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SEQ ID NO:
LENGTH: 0
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SOFTWARE: Fast:
SEQ ID NO 12733
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TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS
FILE REFERENCE: CL001307
LENGTH: 178883
TYPE: DNA
ORGANISM: Human
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Local Similarity 47.8%;
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US-09-949-016-13039/c
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Sequence 142991, Application U
Carent No. 6812339
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                                                                                                Sequence 14 Patent No.
                                                                      GENERAL INFORMATION:
APPLICANT: VENTER,
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CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
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                                   CANT: VENTER, J. Craig et al.

OF INVENTION: POLYMORPHISMS IN KNOWN

OF INVENTION: WITH HUMAN DISEASE, ME

REFERENCE: CL001307
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APPLICATION NUMBER: 60/241,
FILING DATE: 2000-10-20
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NO. 6812339
INFORMATION:
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TINVENTION: WITH HUMAN DISEASE, METHODS
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LENGTH: 1141
TYPE: DNA
ORGANISM: Artificial
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PRIOR FILING DATE: 1999-08-04
NUMBER OF SEQ ID NOS: 23
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NAME/KEY: promoter
NAME/KEY: promoter
LOCATION: (1)..(1141)
OTHER INFORMATION: consensus sequence
09-806-708B-22
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OR APPLICATION NUMBER: 60/237,768
OR FILING DATE: 2000-10-03
OR APPLICATION NUMBER: 60/231,498
OR FILING DATE: 2000-09-08
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RAKYWGWNRABVNSTCTTWKSKTTKVRTSCWANNCRAGDANKDHKWWKWSAAMGVYWNNN
                       GAGCGCATCGCTGGGATTTATGAAAGGCACGAAACTTGACCAATTTGCTAACCAGTGGGC
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CURRENT APPLICATION NUMBER: US/09/9
CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,75
PRIOR APPLICATION NUMBER: 60/237,76
PRIOR APPLICATION NUMBER: 60/237,76
PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,490
PRIOR FILING DATE: 2000-09-08
NUMBER OF SEQ ID NOS: 207017
SOFTWARE. F.
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Matches 52
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Patent No
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APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS
TITLE OF INVENTION: WITH HUMAN DI
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                                                                                    GENERAL INFO
        APPLICANT: Chumakov, Ilya
TITLE OF INVENTION: Polymorphic
FILE REFERENCE: GEN-T112XC1
CURRENT APPLICATION NUMBER: US/C
CURRENT FILING DATE: 1999-06-04
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                                                                         APPLICANT:
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LOCATION: (1)...(265038)
OTHER INFORMATION: n = A
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Chumakov, Ilya
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LOCATION: 92749..97155
OTHER INFORMATION: exon9
FEATURE:
NAME/KEY: exon
LOCATION: 92749..92883
OTHER INFORMATION: exon9bis
FEATURE:
NAME/KEY: exon
LOCATION: 95821..97155
OTHER INFORMATION: exon9ter
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NAME/KEY: exon
LOCATION: 68648..68741
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NAME/KEY: misc_feature
LOCATION: 97156..106746
OTHER INFORMATION: 3'regulation
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NAME/KEY: misc feature
LOCATION: 66647..68647
OTHER INFORMATION: promoter
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CATION: 89484..89649
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R APPLICATION NUMBER: 60/102,324
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ER INFORMATION: exon7
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ATION: 85298..85417
ER INFORMATION: exon4
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INVENTION: Polymorphic Markers

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REFERENCE: GEN-T112XC1 ENT APPLICATION NUMBER: US/0 ENT FILING DATE: 1999-06-04

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PRIOR FILING DATE: 1998-09-28
NUMBER OF SEQ ID NOS: 22
SOFTWARE: Patentin version 3.1
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NAME/KEY: exon
LOCATION: 87496..87522
OTHER INFORMATION: exo
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LOCATION: 86389..8
OTHER INFORMATION:
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LOCATION: 92749..92883
OTHER INFORMATION: exon9bis
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OTHER INFORMATION: exon9ter FEATURE: NAME/KEY: misc_feature LOCATION: 70647.70794
OTHER INFORMATION: homology w.FEATURE: NAME/KEY: misc_feature LOCATION: 68648..68741
OTHER INFORMATION: homology w.FEATURE: NAME/KEY: misc_feature LOCATION: 82205.82229
OTHER INFORMATION: homology w.FEATURE: NAME/KEY: allele LOCATION: 278
OTHER INFORMATION: 99-1601-27.
FEATURE: NAME/KEY: allele LOCATION: 402
OTHER INFORMATION: 99-1601-27.
FEATURE: NAME/KEY: allele LOCATION: 472
OTHER INFORMATION: 99-1601-47.
FEATURE: NAME/KEY: allele LOCATION: 17593
OTHER INFORMATION: 99-13798-2
FEATURE: NAME/KEY: allele LOCATION: 17593
OTHER INFORMATION: 99-13798-2
FEATURE: NAME/KEY: allele LOCATION: 28964
OTHER INFORMATION: 99-13798-2
FEATURE: NAME/KEY: allele LOCATION: 29003
OTHER INFORMATION: 99-13794-1
FEATURE: NAME/KEY: allele LOCATION: 31077
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OTHER INFORMATION: 99-13794-1
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FEATURE: NAME/KEY: allele LOCATION: 34791
OTHER INFORMATION: 99-13812-3
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OTHER INFORMATION: 99-13805-3
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                                Patent No. 6569662
GENERAL INFORMATION:
APPLICANT: Tang, Y.
APPLICANT: Liu, Che
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Best Local
Matches
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       APPLICANT:
APPLICANT:
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NAME/KEY: allele
LOCATION: 49921
OTHER INFORMATION:
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LOCATION: 64239
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APPLICANT: John Tillinghast
APPLICANT: Drmanac, Radoje T.
TITLE OF INVENTION: No. 6569662el Nucleic Ac
TITLE OF INVENTION: Polypeptides
FILE REFERENCE: 784CIP2B
CURRENT APPLICATION NUMBER: US/09/620,312D
CURRENT FILING DATE: 2000-07-19
PRIOR APPLICATION NUMBER: 09/552,317
PRIOR APPLICATION NUMBER: 09/488,725
PRIOR APPLICATION NUMBER: 09/488,725
PRIOR FILING DATE: 2000-01-21
NUMBER OF SEQ ID NOS: 1105
SOFTWARE: pt_FL_genes Version 1.0
SEQ ID NO 351
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                         COMPUTER: IBM Compacione OPERATING SYSTEM: DOS SOFTWARE: FASTSEQ Version 1.5 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/678,3
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TYPE: DNA
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APPLICANT: Xiao, Jian-ping
APPLICANT: Maunus, Robert B.
TITLE OF INVENTION: METHOD FOR CLONING
TITLE OF INVENTION: PRODUCING THE SAPI
TITLE OF INVENTION: ENDONUCLESE
                                                                                                                            S
                                                                                                                                                                                              NUMBER OF SEQUENCES: 8
CORRESPONDENCE ADDRESS:
ADDRESSEE: NEW ENGLAND
STREET: 32 TOZER ROAD
                                                                           COUNTR:
ZIP: 01915
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
MEDIUM TYPE: IBM Compatible
TYPE: IBM COMPATIBLE
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78-304-2
                                                                                                                                                                   CITY:
STATE:
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FILING DATE:
CLASSIFICATION: 435
TORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2.3%; 4.3%; Similarity 57.3%; 59; Conservative
                                                                                                                                                                                                                                                                                                                                                   , Application US/08678304
5663067
                                                                                                                                                                                   BEVERLY
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Xue, Aidong J.
Yang, Yonghong
Wang, Jian-Rui
Zhou, Ping
Ma, Yunging
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                                         US/08/678,304
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Pred. No. 14;
0; Mismatches
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RESTRICTION
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97

NAME: WILLIAMS, REGISTRATION NUMI

NUMBER:

0901

GREGORY

U

NUMBER:

NEB-123

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CURRENT APPLICATION NUMBER: US/09/641,638
CURRENT FILING DATE: 2000-08-16
PRIOR APPLICATION NUMBER: US/09/502,330
PRIOR FILING DATE: 2000-02-11
PRIOR PILING DATE: 2000-02-11
PRIOR PILING DATE: 1999-05-07
PRIOR FILING DATE: 1999-05-07
PRIOR APPLICATION NUMBER: US 09/275,267
PRIOR APPLICATION NUMBER: US 09/275,267
PRIOR APPLICATION NUMBER: US 60/119,917
PRIOR FILING DATE: 1999-03-23
PRIOR FILING DATE: 1999-02-12
NUMBER OF SEQ ID NOS: 1304
SOFTWARE: Patent.pm
SEQ ID NO 232
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US-09-641-638-
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Best Local
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                        OTHER INFO
                                                                    NAME/KEY: misc_binding
LOCATION: 178..197
OTHER INFORMATION: 10-
                                     NAME/KEY: misc_binding LOCATION: 199..218 OTHER INFORMATION: 10-
                                                                                                             LOCATION: 198
OTHER INFORMATION:
                                                                                                                                      NAME/KEY:
                                                                                                                                                             TYPE: DNA
ORGANISM: Homo
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TELEPAX: 508-927-1705
                                                                                                                                                    FEATURE:
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             primer_
1..19
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198
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Chumakov, Ilya
Cohen, Annick
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ilarity 54.7%;
Conservative
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1...1299
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Pred.
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                                                                                                               polymorphic
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TITLE OF INVENTION: BIALLELIC MARKERS DERIVED FROM GENOMIC REGIONS CAR.

TITLE OF INVENTION: GENES INVOLVED IN ARACHIDONIC ACID METABOLISM

FILE REFERENCE: GEN-T114XC2D1

CURRENT APPLICATION NUMBER: US/10/170,097

CURRENT FILING DATE: 2002-06-10

PRIOR APPLICATION NUMBER: US 09/641,638

PRIOR FILING DATE: 2000-08-16

PRIOR PILING DATE: 2000-02-11

PRIOR APPLICATION NUMBER: US 09/502,330

PRIOR APPLICATION NUMBER: US 60/133,200

PRIOR FILING DATE: 1999-03-03-03

PRIOR APPLICATION NUMBER: US 60/139,917

PRIOR PILING DATE: 1999-03-23

PRIOR PILING DATE: 1999-02-12
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Best Lo
Matcher
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Patent
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APPLICANT: Bougueleret
APPLICANT: Chumakov, I
APPLICANT: Cohen, Annio
TITLE OF INVENTION: BIAL
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LOCATION: 403..420
OTHER INFORMATION: downstream
NAME/KEY: misc_binding
LOCATION: 186..210
OTHER INFORMATION: 10-94-198 p
NAME/KEY: misc_feature
LOCATION: 351
                                                                                                                                                                                                                                                                                                                                               SEQ
                                                FEATURE:
NAME/KEY: misc_binding
LOCATION: 199..218
OTHER INFORMATION: 10-94
FEATURE:
NAME/KEY: primer_bind
TOCATION: 1..19
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NUMBER OF SEQ ID NOS: 1304
SOFTWARE: Patent.pm
EQ ID NO 232
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9-641-638-232
                                                                                                                                                                                                                                     FEATURE:
NAME/KEY: allele
LOCATION: 198
OTHER INFORMATION: 10-94-198:
                                                                                                                                                                                                                                                                                                 TYPE: DNA ORGANISM: Homo Sapiens
                                                                                                                                                                                     FEATURE:
NAME/KEY: misc binding
LOCATION: 178..197
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NAM
                                            LOCATION: 1..19 OTHER INFORMATION: upstream
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              Æ/KEY:
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                                                                                                                                                                         ER INFORMATION: 10-94-198.mis1,
  ATION:
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Similarity 54.4%;
52; Conservative
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Chumakov, Ilya
Cohen, Annick
primer_bind 403..420
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Pred. No.
1; Mismatc
                                               amplification
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RESULT 15
US-09-621-976-2446
; Sequence 2446, Application US/09621976
; Patent No. 6639063
; GENERAL INFORMATION:
APPLICANT: Dumas Milne Edwards, J.B.
; APPLICANT: Jobert, S.
; APPLICANT: Giordano, J.Y.
TITLE OF INVENTION: ESTs and Encoded Human FILLE REFERENCE: GENSET.054PR2
CURRENT APPLICATION NUMBER: US/09/621,976
; CURRENT FILING DATE: 2000-07-21
; NUMBER OF SEQ ID NOS: 19335
; SOFTWARE: Patent.pm
; SEQ ID NO 2446
; LENGTH: 430
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Search completed: March Job time : 190 secs
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Best Local S
Matches 53
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Best Local S
Matches 62
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ORGANISM: Homo Bap
FEATURE:
NAME/KEY: CDS
LOCATION: 63..254
1-09-621-976-2446
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NAME/KEY: misc_feature
LOCATION: 351
OTHER INFORMATION: n=a,
10-170-097-232
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FEATURE:
NAME/KEY: misc binding
LOCATION: 186..210
OTHER INFORMATION: 10-9
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Similarity 54.4%;
62; Conservative
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RESULT 1
US-10-527-771-9
Sequence 9, Application US/10527771
Publication No. US20050271683A1
GENERAL INFORMATION:
APPLICANT: University Gent
TITLE OF INVENTION: Ostertagia vaccine
FILE REFERENCE: 2002-015
CURRENT APPLICATION NUMBER: US/10/527,771
CURRENT FILING DATE: 2005-03-11
PRIOR APPLICATION NUMBER: US 10/243,319
PRIOR FILING DATE: 2002-09-13
NUMBER OF SEQ ID NOS: 27
SOFTWARE: Patentin version 3.2
SEQ ID NO 9
LENGTH: 763
TYPE: DNA
ORGANISM: Ostertagia ostertagi
FEATURE:
NAME/KEY: CDS
LOCATION: (11)..(706)
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TITLE OF INVENTION: Ostertagia vaccine
FILE REFERENCE: 2002-015
CURRENT APPLICATION NUMBER: US/10/527,771
CURRENT FILING DATE: 2005-03-11
PRIOR APPLICATION NUMBER: US 10/243,319
PRIOR FILING DATE: 2002-09-13
NUMBER OF SEQ ID NOS: 27
SOFTWARE: Patentin version 3.2
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Publication No. US20050266459A1
GENERAL INFORMATION:
APPLICANT: POULSEN, TIM S.
APPLICANT: NIELSEN, KIRSTEN V.
APPLICANT: NIELSEN, KIRSTEN V.
TITLE OF INVENTION: NUCLEIC ACID PROBES AND NUCLEIC ACID ANALOG PROBES
FILE REFERENCE: 09138.6000-00000
CURRENT APPLICATION NUMBER: US/11/121,086
CURRENT FILING DATE: 2005-05-04
PRIOR APPLICATION NUMBER: 60/567,570
PRIOR FILING DATE: 2004-05-04
NUMBER OF SEQ ID NOS: 107
SOFTWARE: PatentIn version 3.3
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APPLICANT: Wang, David G.
TITLE OF INVENTION: Identification and Mapping of S
TITLE OF INVENTION: Nucleotide Polymorphisms in th
FILE REFERENCE: 108827.135
CURRENT APPLICATION NUMBER: US/09/925,065A
CURRENT FILING DATE: 2001-08-08
PRIOR APPLICATION NUMBER: US 60/243,096
PRIOR APPLICATION NUMBER: US 60/252,147
PRIOR APPLICATION NUMBER: US 60/252,147
PRIOR APPLICATION NUMBER: US 60/250,092
PRIOR APPLICATION NUMBER: US 60/250,092
PRIOR FILING DATE: 2000-11-30
PRIOR FILING DATE: 2000-11-30
PRIOR FILING DATE: 2001-01-16
PRIOR FILING DATE: 2001-01-16
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US-09-929
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US-09-925-065A-611636/c
; Sequence 611636, Application US/09925065A
; Publication No. US20040181048A1
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Publication No.
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CURRENT APPLICATION NUMBER: US/09/9
CURRENT FILING DATE:
                                                NUMBER OF
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LENGTH:
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OR APPLICATION NUMBER: US 60/243,096

OR FILING DATE: 2000-10-24

OR APPLICATION NUMBER: US 60/252,147

OR APPLICATION NUMBER: US 60/250,092

OR APPLICATION NUMBER: US 60/250,092

OR FILING DATE: 2000-11-30

OR APPLICATION NUMBER: US 60/261,766

OR APPLICATION NUMBER: US 60/289,846

OR APPLICATION NUMBER: US 60/289,846

OR APPLICATION NUMBER: US 60/289,846

OR FILING DATE: 2001-05-09
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on No. US20040181048A1
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                             CATION NUMBER: US 6 DATE: 2001-05-09 Q ID NOS: 957086 Mindows
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GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Wang, David G.
APPLICANT: Wang, David G.
TITLE OF INVENTION: Identification and Mapping
TITLE OF INVENTION: Nucleotide Polymorphisms f
                                                                                                                                                                 GENERAL INFORMATION:
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Sequence 524748, Application US/09925065A
Publication No. US20040181048A1
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LENGTH: 538
TYPE: DNA
ORGANISM: Homo sapiens
-09-925-065A-524748
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                                                    APPLICANT: Wang, David G.
TITLE OF INVENTION: Identification
TITLE OF INVENTION: Nucleotide Po-
FILE REFERENCE: 108827.135
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SOFTWARE: FastSEQ for
EQ ID NO 524748
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RENT APPLICATION NUMBER: US/09/925,065A

RENT FILING DATE: 2001-08-08

[OR APPLICATION NUMBER: US 60/243,096

[OR FILING DATE: 2000-10-24

[OR APPLICATION NUMBER: US 60/252,147

[OR FILING DATE: 2000-11-20

[OR FILING DATE: 2000-11-30

[OR APPLICATION NUMBER: US 60/261,766

[OR APPLICATION NUMBER: US 60/261,766
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OF INVENTION: Nucleotide Polymorphii REFERENCE: 108827.135
INT APPLICATION NUMBER: US/09/925,065A
INT FILING DATE: 2001-08-08
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APPLICATION NUMBER: US
FILING DATE: 2001-05-09
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TITLE OF INVENTION: Identification and Mappir
TITLE OF INVENTION: Nucleotide Polymorphisms
FILE REFERENCE: 108827.135
CURRENT APPLICATION NUMBER: US/09/925,065A
CURRENT FILING DATE: 2001-08-08
PRIOR APPLICATION NUMBER: US 60/243,096
PRIOR APPLICATION NUMBER: US 60/252,147
PRIOR APPLICATION NUMBER: US 60/252,147
PRIOR APPLICATION NUMBER: US 60/250,092
PRIOR APPLICATION NUMBER: US 60/261,766
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Sequence 625827, Application US
Publication No. US20040181048A1
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SOFTWARE: FastSEQ for
SEQ ID NO 625826
LENGTH: 570
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PRIOR APPLICATION NUMBER: US (
PRIOR FILING DATE: 2001-05-09
NUMBER OF SEQ ID NOS: 957086
                                                                 LENGTH: 570
TYPE: DNA
ORGANISM: HO
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R FILING DATE: 2000-11-30
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APPLICATION NUMBER: US
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; Sequence 13272, Application US/1099561
; Publication No. US20050272054A1
; GENERAL INFORMATION:
; APPLICANT: CARGILL, Michele et al.
; TITLE OF INVENTION: GENETIC POLYMORPHISMS ASSOCIATED WITH
; TITLE OF INVENTION: CARDIOVASCULAR DISORDERS AND DRUG RESPONTITLE OF INVENTION: DETECTION AND USES THEREOF
; FILE REFERENCE: CL001559
; CURRENT APPLICATION NUMBER: US/10/995,561
; CURRENT FILING DATE: 2004-11-24
; NUMBER OF SEQ ID NOS: 85702
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 13272
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APPLICANT: Wang, David G.
TITLE OF INVENTION: Identification and Mapping of S
TITLE OF INVENTION: Nucleotide Polymorphisms in th
FILE REFERENCE: 108827.135
CURRENT APPLICATION NUMBER: US/09/925,065A
CURRENT FILING DATE: 2001-08-08
PRIOR APPLICATION NUMBER: US 60/243,096
PRIOR FILING DATE: 2000-10-24
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PRIOR APPLICATION NUMBER: US 60/250,092
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PRIOR APPLICATION NUMBER: US 60/261,766
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PRIOR APPLICATION NUMBER: US 60/289,846
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US-09-925-065A-301865
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TYPE: DNA
ORGANISM: Homo
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NAME/KEY: misc_feature
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Pred. No. 1.1e
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CURRENT FILING DATE: 2001-08-08

PRIOR APPLICATION NUMBER: US 60/243,096

PRIOR APPLICATION NUMBER: US 60/252,147

PRIOR FILING DATE: 2000-11-20

PRIOR APPLICATION NUMBER: US 60/250,092

PRIOR FILING DATE: 2000-11-30

PRIOR APPLICATION NUMBER: US 60/261,766

PRIOR APPLICATION NUMBER: US 60/261,766
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SOFTWARE: Pascon
SEQ ID NO 301865
FONGTH: 574
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Best Local &
Matches 93
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SOFTWARE: FastSEQ for W
SEQ ID NO 625825
LENGTH: 570
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-09-925-065A-625825/c
Sequence 625825, Application US/09925065A
Publication No. US20040181048A1
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Matches
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                                                                                                                                                                                                                                                                                                                  FILING DATE: 2001-01-16
APPLICATION NUMBER: US
FILING DATE: 2001-05-09
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OF INVENTION: Identification

OF INVENTION: Nucleotide Poly

PEFERENCE: 108827.135
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Similarity 52.1%;
74; Conservative
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Similarity 47.7%;
3; Conservative
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A-625
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SEQ ID NOS: 957086
PastSEQ for Windows
ACCTGTGTGACACGC
                                   GAACCCGGGAGGCGGAAGCTGCAGTGAGCTGAGATTGTGCCACTGCATTCCAGCCCAGGC
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RESULT
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                                                      APPLICANT: Wang, David G.
TITLE OF INVENTION: Identification and Mapping of
TITLE OF INVENTION: Nuclectide Polymorphisms in
FILE REFERENCE: 108827.135
CURRENT APPLICATION NUMBER: US/09/925,065A
CURRENT FILING DATE: 2001-08-08
PRIOR APPLICATION NUMBER: US 60/243,096
PRIOR FILING DATE: 2000-10-24
PRIOR APPLICATION NUMBER: US 60/252,147
PRIOR APPLICATION NUMBER: US 60/252,147
PRIOR APPLICATION NUMBER: US 60/250,092
PRIOR APPLICATION NUMBER: US 60/250,092
PRIOR APPLICATION NUMBER: US 60/261,766
PRIOR APPLICATION NUMBER: US 60/261,766
PRIOR FILING DATE: 2001-01-16
PRIOR FILING DATE: 2001-05-09
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LENGTH: 576
TYPE: DNA
ORGANISM: Homo sapiens
JS-09-925-065A-193665
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CURRENT FILING DATE: 2001-08-08
PRIOR APPLICATION NUMBER: US 60/243,096
PRIOR FILING DATE: 2000-10-24
PRIOR APPLICATION NUMBER: US 60/252,147
PRIOR FILING DATE: 2000-11-20
PRIOR APPLICATION NUMBER: US 60/250,092
PRIOR FILING DATE: 2000-11-30
PRIOR APPLICATION NUMBER: US 60/261,766
PRIOR APPLICATION NUMBER: US 60/261,766
PRIOR APPLICATION NUMBER: US 60/289,846
PRIOR FILING DATE: 2001-05-09
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   NUMBER OF SOFTWARE:
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TITLE OF INVENTION: Identification and Mapping of Sir
TITLE OF INVENTION: Nucleotide Polymorphisms in the
FILE REFERENCE: 108827.135
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OR APPLICATION NUMBER: US 60/261,766
OR FILING DATE: 2001-01-16
OR APPLICATION NUMBER: US 60/289,846
OR FILING DATE: 2001-05-09
BER OF SEQ ID NOS: 957086
IWARE: FastSEQ for Windows Version 4.0
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25-065A-745205
ince 745205, Application US
cation No. US20040181048A1
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)nce 193665, Application US/09925065A
.cation No. US20040181048A1
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l Similarity 49.7%;
83; Conservative (
SEQ ID NOS:
FastSEQ for
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                       Version
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Pred. No. 1
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                                            RESULT
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CURRENT APPLICATION NUMBER: US/09/925,
CURRENT FILING DATE: 2001-08-08
PRIOR APPLICATION NUMBER: US 60/243,09
PRIOR APPLICATION NUMBER: US 60/252,14
PRIOR APPLICATION NUMBER: US 60/252,14
PRIOR FILING DATE: 2000-11-20
PRIOR APPLICATION NUMBER: US 60/250,09
PRIOR FILING DATE: 2000-11-30
PRIOR APPLICATION NUMBER: US 60/261,76
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Best Loc
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Best Local
Matches
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-09-925-065A-189082/c
Sequence 189082, Application US
Publication No. US20040181048A1
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SULT 15;-09-925-065A-250352/c;-09-925-065A-250352/c
Sequence 250352, Application US
Publication No. US20040181048A1
GENERAL INFORMATION:
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SOFTWARE: FastSEQ for
SEQ ID NO 189082
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TITLE OF IN
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TYPE: DNA
ORGANISM:
FEATURE:
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ORGANISM: Ho
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LOCATION: 501
OTHER INFORMATION: n =
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APPLICATION NUMBER: US
FILING DATE: 2001-05-09
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OF INVENTION:
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11 Similarity 51.4%;
72; Conservative
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OF INVENTION: Identification and Mapping of Single

OF INVENTION: Nucleotide Polymorphisms in the Human Genome

REFERENCE: 108827.135
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ALIGNMENTS

RESULT US-10-2

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US-10-243-319C-9
; Sequence 9, Application US/10243319C
; Publication No. US20040052817A1
; GENERAL INFORMATION:
APPLICANT: Universiteit Gent
TITLE OF INVENTION: Ostertagia vaccine
FILE REFERENCE: UniGent Ostertagia
CURRENT APPLICATION NUMBER: US/10/243,319
CURRENT FILING DATE: 2002-09-13
NUMBER OF SEQ ID NOS: 14
; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 9
; LENGTH: 763
TYPE: DNA
ORGANISM: Ostertagia ostertagi
FEATURE:
NAME/KEY: CDS
; LOCATION: (11)...(706)
US-10-243-319C-9
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TITLE OF INVENTION: Ostertagia vaccine
FILE REFERENCE: UniGent Ostertagia
CURRENT APPLICATION NUMBER: US/10/243,31
CURRENT FILING DATE: 2002-09-13
NUMBER OF SEQ ID NOS: 14
SOFTWARE: Patentin version 3.2
SEQ ID NO 7
LENGTH: 828
                                                                                                                                                                                                                                                                          Sequence 7, Application US/1024331
Publication No. US20040052817A1
GENERAL INFORMATION:
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Best Local S
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                                                                                                                                                          TYPE: DNA
ORGANISM: Ostertagia (
FEATURE:
NAME/KEY: CDS
LOCATION: (11)..(721)
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63.6%;
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Sequence 84, Application US/10388838; Publication No. US20040180344A1
GENERAL INFORMATION:
APPLICANT: David W. Morris
APPLICANT: Marc Malandro
TITLE OF INVENTION: Novel Therapeutic Target
FILE REFERENCE: 529452001600
CURRENT APPLICATION NUMBER: US/10/388,838
CURRENT FILING DATE: 2003-03-14
NUMBER OF SEQ ID NOS: 114
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 84
LENGTH: 260803
TYPE: DNA
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NAME/KEY: misc_feature LOCATION: (1)...(260803)
OTHER INFORMATION: n = 1-10-388-838-84
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INFORMATION: n = A
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Similarity 45.7%;
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ATGCAATGATCCCCTACCGATAAATACCAGCCTGGCTCAAAATATCGCTAGATGGCTGTA
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RESULT 5
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; Sequence 5625, Application US/10741601
; Publication No. US20040166519A1
; GENERAL INFORMATION:
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PRIOR APPLICATION NUMBER: US 60
PRIOR APPLICATION NUMBER: US 60
PRIOR APPLICATION NUMBER: US 60
PRIOR FILING DATE: 2001-01-16
PRIOR APPLICATION NUMBER: US 60
PRIOR FILING DATE: 2001-05-09
NUMBER OF SEQ ID NOS: 957086
SOFTWARE: FastSEQ for Windows V
                                                                                                     TITLE OF INVENTION: STENOSIS FILE REFERENCE: CL001500 CURRENT APPLICATION NUMBER: CCURRENT FILING DATE: 2003-12 NUMBER OF SEQ ID NOS: 26415 SOFTWARE: FastSEQ for Windows SEQ ID NO 5625
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  Query Match
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Matches 100; Conser
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                                                                                                                                                                                  TITLE
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                                                                              TYPE: DNA
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LENGTH:
TYPE: DN
                                                                    ORGANISM:
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TLE OF INVENTION: Identification and Mapping of
TLE OF INVENTION: Nucleotide Polymorphisms in
LE REFERENCE: 108827.135
LE REFERENCE: 108827.135
RRENT APPLICATION NUMBER: US/09/925,065A
RRENT FILING DATE: 2001-08-08
IOR APPLICATION NUMBER: US 60/243,096
IOR FILING DATE: 2000-10-24
IOR APPLICATION NUMBER: US 60/252,147
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US-10-367-094-122/c; Sequence 122, Application US/10367094; Publication No. US20040170982A1

GENERAL INFORMATION:

APPLICANT: David W. Morri

Morris

Therapeutic

Targets

İn

Cancer

APPLICANT: Marc Malauter TITLE OF INVENTION: Novel Ther FILE REFERENCE: 529452001500 CURRENT APPLICATION NUMBER: US CURRENT FILING DATE: 2003-02-NUMBER OF SEQ ID NOS: 203

SEQ

LENGTH:

421609

SOFTWARE:

Windows

Version 4.0

2003-02-

US/10/367,094

RESULT

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CURRENT APPLICATION NUMBER: US/10/087,192; CURRENT FILING DATE: 2002-03-01; PRIOR APPLICATION NUMBER: US 09/747,377; PRIOR FILING DATE: 2000-12-22; PRIOR APPLICATION NUMBER: US 09/798,586; PRIOR FILING DATE: 2001-03-02; NUMBER OF SEQ ID NOS: 2059; SOPTWARE: FastSEQ for Windows Version 4.0; SEQ ID NO 826; LENGTH: 194883
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TITLE OF INVENTION: NOVEL COMPOSITIONS AND
TITLE OF INVENTION: CANCER
FILE REFERENCE: 529452000122
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ORGANI
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187-192-826/c
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Pred. No. 85;
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RESULT 9
US-10-363-345A-10700
; Sequence 10700, Application US/10
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· GENERAL INFORMATION:
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US-10-363-345A-10699/c
; Sequence 10699, Application US/
; Publication No. US20040234960A1
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ORGANISM: Mus musculus
FEATURE:
NAME/KEY: misc_feature
LOCATION: (1)...(421609)
OTHER INFORMATION: n = A,
-10-367-094-122
APPLICANT:
APPLICANT:
APPLICANT:
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APPLICANT: Christian Pie
APPLICANT: Kurt Berlin
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Similarity 50.3%;
85; Conservative
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GENERAL INFORMATION:

APPLICANT: Alexander Olek

APPLICANT: Christian Piepenbrock

APPLICANT: Kurt Berlin

TITLE OF INVENTION: Diagnosis of illnesses or

TITLE OF INVENTION: illnesses

FILE REFERENCE: 82011

CURRENT APPLICATION NUMBER: US/10/363,483A

CURRENT FILING DATE: 2003-03-03

NUMBER OF SEQ ID NOS: 40712

SEQ ID NO 10699

LENGTH: 1003

TYPE: DNA
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CURRENT APPLICATION NUMBER: US/10/363,345A
CURRENT FILING DATE: 2003-03-03
NUMBER OF SEQ ID NOS: 40712
SEQ ID NO 10700
LENGTH: 1003
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OF INVENTION: Identification and Mapping of INVENTION: Polymorphisms in the Human REFERENCE: 108827.129

ENT APPLICATION NUMBER: US/10/027,632
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APPLICATION NUMBER:
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APPLICANT: Zhou, Yihua
APPLICANT: Cao, Yongwei
APPLICANT: Wu, Wei
APPLICANT: Boukharov, Andrey A.
APPLICANT: Boukharov, Andrey A.
APPLICANT: Li, Ping
TITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules Assumith of Invention: Plants and Uses Thereof for Plant Improvement
FILLS OF INVENTION : Plants and Uses Thereof for Plant Improvement
FILLS REFERENCE: 38-21(53221)B
CURRENT APPLICATION NUMBER: US/10/437,963
CURRENT FILING DATE: 2003-05-14
NUMBER OF SEQ ID NOS: 204966
SEQ ID NO 51217
LENGTH: 388
TYPE: PN"
                                                                                                                                                                     ; LENGTH: 653458
; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: (1)...(653458)
; OTHER INFORMATION: n = A,7
US-10-461-862-4
                                                                                                                                                                                                                                                                                   Sequence 4, Application US/10461862; Publication No. US20050090434A1; GENERAL INFORMATION:
APPLICANT: David W. Morris
APPLICANT: Marc S. Malandro
TITLE OF INVENTION: Novel Therapeutic Ta
FILE REFERENCE: 529452001800
CURRENT APPLICATION NUMBER: US/10/461,86
CURRENT FILING DATE: 2003-06-13
NUMBER OF SEQ ID NOS: 184
SOFTWARE: FastSEQ for Windows Version 4.
SEQ ID NO 4
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ORGANISM: Oryza sativa
FEATURE:
OTHER INFORMATION: Clone
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ALIGNMENTS

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/organism="Telacorbagia circumcincta" /mol_type="mRNA" /db_xref="taxon:45464" /clone="Tc_ad2_28F03" /sex="mixed") - -	High quality sequence stop: 519. Location/Oualifiers	Plate: 28 row: F column: 03	BACKWARD: TFXPCRF1	Sequencing was performed by the Pathogen Sequencing Unit, Sanger Centre, Cambridge, UK (Neil Hall, Mike Quail & Bart Barrell).	performed by Ye Jieru, ICAPB, University of	Moredun Research Institute, Midlothian, UK.	ll: mark.blaxter@ed.ac.uk	Fax: +44 131 670 5450	+44 131 650	rane, will a partarilla, near marie woar, earimaryii,	gn Buildings West Mains Doad		Unpublished (2000)	Edinburgh University/Sanger Centre Nematode EST Project	M.L., Parkinson, J., W	0 766)	idae; Ostertagiinae; Telac	m	Teladorgagia circumcincta		CB036994.1 GI:27756239		ט טי	gia circumcincta adults l	766 by mony linear	

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/clone_lib="Teladorsagia circumcincta adults library 2
/note="Vector: pTriplEx2; Site_1: BamHI; Site_2: BamHI
Teladorsagia circumcincta is a parasitic nematode. The
library was constructed from mRNA from Teladorsagia
circumcincta mixed adults."
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Institute of Cell, Animal and Population Biology
University of Edinburgh
Ashworth Labs, King's Buildings, West Mains Road, Edin
3JT, UK.
Tel: +44 131 650 6760
Fax: +44 131 670 5450
Email: mark.blaxter@ed.ac.uk
The library was prepared by Dr Diane Redmond and Dr Day
Moredun Research Institute, Midlothian, UK.
PCR was performed by Ye Jieru, ICAPB, University of Edentre, Cambridge, UK (Neil Hall, Mike Quail & Bart Bart Bart PCR PRimers
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Plate: 41 row: E column: 11
Seq primer: TEXF1
High quality sequence start: 5
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Teladorsagia circumcincta
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Eukaryota; Metazoa; Nematoda; Chromadorea; Rhabditida; Strongylida;
Trichostrongyloidea; Haemonchidae; Ostertagiinae; Teladorsagia.
1 (bases 1 to 757)
Blaxter, M.L., Parkinson, J., Whitton, C., Daub, J., Guiliano, D.,
Hall, N., Quayle, M. and Barrell, B.
Edinburgh University/Sanger Centre Nematode EST Project
Unpublished (2000)
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78; Conservative
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/mol_type="mRNA"
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/db_xref="taxon:45464"
/clone="Tc_ad2_41E11"
/dev_stage="adult"
/clone_lib="Teladorsagia circumcincta adults library
/note="Vector: pTriplex2; Site_1: BamHI; Site_2: BamH
Teladorsagia circumcincta is a parasitic nematode. Th
library was constructed from mRNA from Teladorsagia
circumcincta mixed adults."
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Fax: +44 131 670 5450

Email: mark.blaxter@ed.ac.uk

The library was prepared by Dr Diane Redmond and Dr Da

Moredun Research Institute, Midlothian, UK.

PCR was performed by Ye Jieru, ICAPB, University of Ed

Sequencing was performed by the Pathogen Sequencing Un

Centre, Cambridge, UK (Neil Hall, Mike Quail & Bart Ba

PCR PRimers
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Blaxter, M.L., Parkinson, J., Whitton, C., D
Hall, N., Quayle, M. and Barrell, B.
Edinburgh University/Sanger Centre Nemato
Unpublished (2000)
Contact: Blaxter ML
Institute of Cell, Animal and Population
University of Edinburgh
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RESULT 4
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/clone_lib="Teladorsagia circumcincta adults librar,
/note="Vector: pTriplEx2; Site_1: BamHI; Site_2: Ban
Teladorsagia circumcincta is a parasitic nematode. '
library was constructed from mRNA from Teladorsagia
circumcincta mixed adults."
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Blaxter,M.L., Parkinson,J., Whitton Hall,N., Quayle,M. and Barrell,B. Edinburgh University/Sanger Centre Unpublished (2000)
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1 (bases 1 to 760)
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Teladorsagia circumcincta
Eukaryota; Metazoa; Nematoda;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Tel: +44 131 650 6760

Fax: +44 131 670 5450

Email: mark.blaxter@ed.ac.uk

The library was prepared by Dr Diane Redmond and Dr

Moredun Research Institute, Midlothian, UK.

PCR was performed by Ye Jieru, ICAPB, University of

Sequencing was performed by the Pathogen Sequencing

Centre, Cambridge, UK (Neil Hall, Mike Quail & Bart

PCR PRimers
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High quality sequence stop: Location/Qualifiers
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3JT, UK.
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                                                 AGTAGCAGCACAACAGATTGCGCCATGCAATGATCCCCTACCGATAAATACCAGCCTGGC
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/db_xref="taxon:45464"
/clone="Tc_ad2_44A12"
/sex="mixed"
/dev_stage="adult"
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Fax: +44 131 670 5450
Email: mark.blaxter@ed.ac.uk
The library was prepared by Dr Diane Redmond and Dr Da.
Moredun Research Institute, Midlothian, UK.
PCR was performed by Ye Jieru, ICAPB, University of Ed.
Sequencing was performed by the Pathogen Sequencing Un.
Centre, Cambridge, UK (Neil Hall, Mike Quail & Bart Ba.
PCR PRimers
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Blaxter, M.L., Parkinson, J., Whitton, C., Daub, J. Hall, N., Quayle, M. and Barrell, B. Edinburgh University/Sanger Centre Nematode ES Unpublished (2000)
Contact: Blaxter ML
Institute of Cell, Animal and Population Biold University of Edinburgh Ashworth Labs, King's Buildings, West Mains Ro
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CB038112
CB038112.1
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3JT, UK.
Tel: +44
Fax: +44
                                                           BACKWARD: T7
Plate: 42 r
Seq primer:
High quality
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Teladorsagia circumcincta
Eukaryota; Metazoa; Nematoda; Chromadorea; Rhabditida; Strong
Trichostrongyloidea; Haemonchidae; Ostertagiinae; Teladorsagi
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RESULT 6
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Best Local Similarity 79.0%;
Matches 579; Conservative
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/sex="mixed"
/dev_stage="adult"
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/note="Vector: pTriplEx2; Site_1: BamHI; Site_2: BamF
Teladorsagia circumcincta is a parasitic nematode. The stary was constructed from mRNA from Teladorsagia circumcincta mixed adults."
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); Mismatches 148;
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Tel: +44 131 650 6760

Fax: +44 131 670 5450

Email: mark.blaxter@ed.ac.uk

The library was prepared by Dr Diane Redmond and Dr Day

Moredun Research Institute, Midlothian, UK.

PCR was performed by Ye Jieru, ICAPB, University of Edine Sequencing was performed by the Pathogen Sequencing University of Edine Sequencing University of Edine PCR PRimers

FORWARD: TEXPCRF1

BACKWARD: T7PL

Plate: 21 row: B column: 12
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Blaxter, M.L., Parkinson, J., Whitton, C., Daub, J., Guilia: Hall, N., Quayle, M. and Barrell, B.
Edinburgh University/Sanger Centre Nematode EST Project Unpublished (2000)
Contact: Blaxter ML
Institute of Cell, Animal and Population Biology
University of Edinburgh
University of Edinburgh
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Teladorsagia circumcincta is a parasitic nematode. The
library was constructed from mRNA from Teladorsagia
circumcincta mixed adults."
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Pred. No. 3.2e-132;
); Mismatches 144;
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Contact: Blaxter ML
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Teladorsagia circumcincta
Eukaryota; Metazoa; Nematoda; Chromadorea; Rhabditida; Strongylida;
Trichostrongyloidea; Haemonchidae; Ostertagiinae; Teladorsagia.
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Tc_ad2_28D02_TEXF1 Teladorsagia
Teladorsagia circumcincta cDNA c
CAD56659 Ancyclostoma-secreted p
                                                                                                                                                                                   Tel: +44 131 650 6760
Fax: +44 131 670 5450
Email: mark.blaxter@ed.ac.uk
The library was prepared by Dr Diane Redmond and
Moredun Research Institute, Midlothian, UK.
PCR was performed by Ye Jieru, ICAPB, University
Sequencing was performed by the Pathogen Sequence
Centre, Cambridge, UK (Neil Hall, Mike Quail & Ba
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University of Edinburgh
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/clone="Tc_ad2_28
/sex="mixed"
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Fax: +44 131 670 5450
Email: mark.blaxter@ed.ac.uk
The library was prepared by Dr Diane Redmond and Dr Da
Moredun Research Institute, Midlothian, UK.
PCR was performed by Ye Jieru, ICAPB, University of Ed
Sequencing was performed by the Pathogen Sequencing Un
Centre, Cambridge, UK (Neil Hall, Mike Quail & Bart Ba
PCR PRimers
FORWARD: TEXPCRF1
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Contact: Blaxter ML
Institute of Cell, Animal and Population Bi University of Edinburgh
Ashworth Labs, King's Buildings, West Mains
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CB038204
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3JT, UK.
Tel: +44 131 650 6760
Fax: +44 131 670 5450
Email: mark.blaxter@ed.ac.uk
                                                                                                                                                                                                                   Unpubliance Contact: Blaxter ML Contact: Blaxter ML Institute of Cell, Animal and University of Edinburgh Coloreth Labs, King's Building
                                                                                         The library was prepared by Dr Diane Redmond and Dr Moredun Research Institute, Midlothian, UK. PCR was performed by Ye Jieru, ICAPB, University of Sequencing was performed by the Pathogen Sequencing Centre, Cambridge, UK (Neil Hall, Mike Quail & Bart PCR PRimers FORWARD: TEXPCRF1
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Edinburgh University/Sanger Centre Nematode EST Unpublished (2000)
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Teladorsagia circumcincta
Teladorsagia circumcincta
Eukaryota; Metazoa; Nematoda; Chromadorea; Rhabditida; Strongylida; Trichostrongyloidea; Haemonchidae; Ostertagiinae; Teladorsagia.

1 (bases 1 to 754)
Blaxter, M.L., Parkinson, J., Whitton, C., Daub, J., Guiliano, D.,
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/mol_type="mRNA"
/db_xref="taxon:45464"
/clone="Tc_ad2_43A10"
/sex="mixed"
/dev_stage="adult"
/clone_lib="Teladorsagia circumcincta adults library 2"
/note="Vector: pTriplEx2; Site_1: BamHI; Site_2: BamHI;
Teladorsagia circumcincta is a parasitic nematode. The
library was constructed from mRNA from Teladorsagia
circumcincta mixed adults."
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Teladorsagia circumcincta cDNA clone Tc ad2 44F10 5' similar to
CAD56659 Ancyclostoma-secreted protein-like prote Ostertagia, mRI
sequence.

VCB038345
CB038345.1 GI:27757590
EST.
Teladorsagia circumcincta
ENTeladorsagia circumcincta
Eukaryota; Metazoa; Nematoda; Chromadorea; Rhabditida; Strongylidelications and to 771)
Trichostrongyloidea; Haemonchidae; Ostertagiinae; Teladorsagia.

I (bases 1 to 771)
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BACKWARD: T7PL
Plate: 44 row: F
Seq primer: TEXF1
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Hall,N., Quayle,M. and Barrell,B.
Edinburgh University/Sanger Centre Nematode EST
Unpublished (2000)
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AGTAGCAGCACAACAGATTGCGCCATGCAATGATCCCCCTACCGATAAATACCAGCCTGGC
                                                                                                                   TATTCGTCGGAACATAGCACTTGAAAAGAGCTTGGTGAACT--
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/clone="Tc_ad2_44F10"
/sex="mixed"
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/mol_type="mRNA"
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                                                                               Tel: +44 131 650 6760
Fax: +44 131 670 5450
Email: mark.blaxter@ed.ac.uk
The library was prepared by Dr Diane Redmond and Dr
Moredun Research Institute, Midlothian, UK.
PCR was performed by Ye Jieru, ICAPB, University of
Sequencing was performed by the Pathogen Sequencing
Centre, Cambridge, UK (Neil Hall, Mike Quail & Bart
PCR PRimers
FORWARD: TEXPCRF1
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Blaxter, M.L., Parkinson, J., Whitton, C. Hall, N., Quayle, M. and Barrell, B.
Edinburgh University/Sanger Centre Nem Unpublished (2000)
Contact: Blaxter ML
Institute of Cell, Animal and Populati University of Edinburgh
Ashworth Labs, King's Buildings, West
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Eukaryota; Metazoa; Haemonchidae; Ostertagiinae; Teladorsagia.
Trichostrongyloidea; Haemonchidae; Ostertagiinae; Teladorsagia.
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/clone_lib="Teladorsagia circumcincta adults library
/note="Vector: pTriplEx2; Site_1: BamHI; Site_2: Ban
Teladorsagia circumcincta is a parasitic nematode. '
library was constructed from mRNA from Teladorsagia
circumcincta mixed adults."
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/mol_type="mRNA"
/db_xref="taxon:45464"
/clone="Tc_ad2_36B07"
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Fax: +44 131 670 5450

Email: mark.blaxter@ed.ac.uk

The library was prepared by Dr Diane Redmond and Dr

Moredun Research Institute, Midlothian, UK.

PCR was performed by Ye Jieru, ICAPB, University of Sequencing was performed by the Pathogen Sequencing Centre, Cambridge, UK (Neil Hall, Mike Quail & Bart PCR PRimers
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/note="Vector: pTriplEx2; Site_1: BamHI; Site_2: BamFieladorsagia circumcincta is a parasitic nematode. The library was constructed from mRNA from Teladorsagia circumcincta mixed adults."
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CB036489
CB036489.1
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Plate: 22
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Teladorsagia circumcincta
Teladorsagia circumcincta
Eukaryota; Metazoa; Nematoda; Chromadorea; Rhabditida; Strongyl
Trichostrongyloidea; Haemonchidae; Ostertagiinae; Teladorsagia.
1 (bases 1 to 739)
Blaxter, M.L., Parkinson, J., Whitton, C., Daub, J., Guiliano, D.,
Hall, N., Quayle, M. and Barrell, B.
Edinburgh University/Sanger Centre Nematode EST Project
Unpublished (2000)
                                                                                                                                                                                      Tel: +44 131 650 6760

Fax: +44 131 670 5450

Email: mark.blaxter@ed.ac.uk

The library was prepared by Dr Diane Redmond and Dr

Moredun Research Institute, Midlothian, UK.

PCR was performed by Ye Jieru, ICAPB, University of

Sequencing was performed by the Pathogen Sequencing

Centre, Cambridge, UK (Neil Hall, Mike Quail & Bart

PCR PRimers
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TC_ad2_22E03_TEXF1 Teladorsagia circumcincta adults library 2
Teladorsagia_circumcincta cDNA clone Tc_ad2_22E03 5' similar
CAD56659 Ancyclostoma-secreted protein-like_prote Ostertagia,
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/mol_type="mRNA"
/db_xref="taxon:45464"
/clone="Tc_ad2_22E03"
/sex="mixed"

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Tc_ad2_39D08_TEXF1 Teladorsagia
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/clone_lib="Teladorsagia circumcincta adults library 2
/note="Vector: pTriplEx2; Site_1: BamHI; Site_2: BamHI
Teladorsagia circumcincta is a parasitic nematode. The
library was constructed from mRNA from Teladorsagia
circumcincta mixed adults."
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The library was prepared by Dr Diane Redmond and Dr
Moredun Research Institute, Midlothian, UK.
PCR was performed by Ye Jieru, ICAPB, University of
Sequencing was performed by the Pathogen Sequencing
Centre, Cambridge, UK (Neil Hall, Mike Quail & Bart
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Institute of Cell, Animal and Population Biology
University of Edinburgh
Ashworth Labs, King's Buildings, West Mains Road
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Edinburgh University/Sanger Centre Nematode EST Unpublished (2000)
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Blaxter, M.L., Parkinson, J., Whitton, C., Daub, J., Guiliano, D.,
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/db_xref="taxon:4546
/clone="Tc_ad2_39D08
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Eukaryota; Metazoa; Nematoda; Chromadorea; Rhabditida; Strongylida
Trichostrongyloidea; Haemonchidae; Ostertagiinae; Teladorsagia.

1 (bases 1 to 758)
Blaxter, M.L., Parkinson, J., Whitton, C., Daub, J., Guiliano, D.,
Hall, N., Quayle, M. and Barrell, B.
Edinburgh University/Sanger Centre Nematode EST Project
Unpublished (2000)
Contact: Blaxter ML
Institute of Cell, Animal and Population Biology
University of Edinburgh
University of Edinburgh
                                                                                                                                                                                   Tel: +44 131 650 6760

Fax: +44 131 670 5450

Email: mark.blaxter@ed.ac.uk

The library was prepared by Dr Diane Redmond and Dr

Moredun Research Institute, Midlothian, UK.

PCR was performed by Ye Jieru, ICAPB, University of

Sequencing was performed by the Pathogen Sequencing

Centre, Cambridge, UK (Neil Hall, Mike Quail & Bart
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/sex="mixed"
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                                    organism="Teladorsagia/mol_type="mRNA"
/db_xref="taxon:45464"
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MEDLINE=96215086; PubMed=8636085; DOI=10.1074/jbc.271.12

Hawdon J.M., Jones B.F., Hoffman D.R., Hotez P.J.;

"Cloning and characterization of Ancylostoma-secreted proprotein associated with the transition to parasitism by hookworm larvae.";

J. Biol. Chem. 271:6672-6678(1996).

-!- FUNCTION: Associated with the transition to parasiting infective hookworm larvae.

-!- SUBCELLULAR LOCATION: Secreted (Potential).
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Mol. Biochem. Parasitol. 126:201-2

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Name=aspl;

Ancylostoma ceylanicum.

Eukaryota; Metazoa; Nematoda; Chromadorea; Rhabditida; St.

Eukaryota; Metazoa; Nematoda; Chromadorea; Rhabditida; St.

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NUCLEOTIDE SEQUENCE.

A Zhan B., Liu Y., Hotez P.J., Hawdon J.;

A Zhan B., Liu Y., Hotez P.J., Hawdon J.;

Submitted (JJL-2002) to the EMBL/GenBank/DDBJ databases.

R EMBL; AY136548; AAN11402.1; -; mRNA.

GO; GO:0005576; C:extracellular region; IEA.

Interpro; IPR002123; N5_allergen.

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R PFANTS; PR00837; V5TPXLIKE.

R PRINTS; PR00837; V5TPXLIKE.

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R SMART; SM00198; SCP; 2.

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4BC8295F5D3035F9 CRC64;
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7153
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EMBL; AF077402; AAD13339.1; -; mRNA.
GO; GO:0005576; C:extracellular region; II
InterPro; IPR001283; Allrgn_V5/Tpx1.
InterPro; IPR002413; V5_allergen.
Pfam; PF00188; SCP; 2.
PRINTS; PR00838; V5ALLERGEN.
PRINTS; PR00837; V5TPXLIKE.
SMART; SM00198; SCP; 2.
PROSITE; PS01009; CRISP_1; 1.
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STRAIN-Shanghai;
MEDLINE-99151774
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077153 981LA
077153;
01-NOV-1998
01-NOV-1998
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                                              SMART; SI
PROSITE;
Signal.
SIGNAL
CHAIN
SEQUENCE
ery Match
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Ancylostomatoidea;
NCBI_TaxID=51022;
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Metazoa; Nematoda; C
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Pred. No. 3.5e-
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G., Zheng
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Score Pred.
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RESULT 6
076744 NECAM
1D 076744 NECAM PRELIMINARY; PRT; 424 AA.
AC 076744;
DT 01-NOV-1998 (TrEMBLrel. 08, Created)
DT 01-NOV-1998 (TrEMBLrel. 24, Last amoutation update)
DT 01-NOV-1998 (TrEMBLrel. 24, Last annotation update)
DT 01-UJUN-2003 (TrEMBLrel. 24, Last annotation update)
DT 01-NOV-1998 (TrEMBLrel. 24, Last annotation update)
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EMBL; AF079521; AAD13340.1; -; mRNA.

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GO; GO:0005576; C:extracellular region; IEA.

InterPro; IPR001283; Allrgn_V5/Tpx1.

InterPro; IPR002413; V5_allergen.

Pfam; PF00188; SCP; 2.

PRINTS; PR00838; V5ALLERGEN.

PRINTS; PR00837; V5TPXLIKE.

SMART; SM00198; SCP; 2.

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"Variation between ASP-1 molecul
and the United States.";

RL J. Parasitol. 86:181-185(2000).
DR EMBL; AF132291; AAD31839.1; -; mR
DR GO; GO:0005576; C:extracellular re
GO; GO:0005576; C:extracellular re
InterPro; IPR001283; Allrgn_V5/Tpx.
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DR PRINTS; PR00838; V5ALLERGEN.
DR PRINTS; PR00837; V5TPXLIKE.
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Q1-NOV-1999 (TrEMBLrel. 12, Last sequence update)
Q1-NOV-2003 (TrEMBLrel. 24, Last annotation update)
Ancylostoma-secreted protein 1 precursor.
Ancylostoma caninum (Dog hookworm).
Eukaryota; Metazoa; Nematoda; Chromadorea; Rhabditida
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Xiao S.-H., Feng
1 ASP-1 molecules
                                         LAVSCIYNGVGYITNQPMWETGQACQTGADCSTYKNSGCEDGLCTKGPDVP
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Pred. No. 6.4e-1
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RESULT 8
QBMQ79 CAREL
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PRT; 424 AA.
AC QBMQ79;
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DT 01-OCT-2002 (TrEMBLrel. 24, Last sequence update)
DT 01-OCT-2003 (TrEMBLrel. 24, Last sequence update)
DT 01-UN-2003 (TrEMBLrel. 24, Last sequence update)
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RAMEDLINE-90-11, ORFWames-F11C7.3;
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HSSP; Q05110; 1QNX.
Ensembl; F11C7.3; Caenorhabditis WormBase; WBGene00006886; vap-1.
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Name=vap-1; ORFNames=F11C7
Caenorhabditis elegans.
Eukaryota; Metazoa; Nematoc
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NCBI_TaxID=6239;
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The C.briggae Sequencing Consortium; Submitted (SEP-2003) to the EMBL/GenBank/DDBJ databases - i - CAUTION: The sequence shown here is derived from an
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CAUTION: The sequence shown here is derived from an
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RESULT 14

P90959 CAREEL

ID P90959;
AC P90959;
DT 01-MAY-1997 (TrEMBLrel. 03, Last sequence update)
DT 01-MAY-1997 (TREMBLrel. 03, Last sequence update)
DT 01-MAY-1997 (TREMBLrel. 24, Last annotation update)
DT 01-JUN-2003 (TREMBLrel. 24, Last annotation update)
DE Hypothetical protein T05A10.4.

GN ORFNames-T05A10.4;
GN Caenorhabditis elegans.
OC Eukaryota; Metazoa; Nematoda; Chromadorea; Rhabditida;
OC Rhabditidae; peloderinae; Caenorhabditis.
OX NCBI_TaxID=6239;
RI [1]
RP NUCLEOTIDE SEQUENCE [LARGE SCALE GENOMIC DNA].
RP NUCLEOTIDE SEQUENCE [LARGE SCALE GENOMIC DNA].
RC STRAIN=Bristol N2;
RC STRAIN=Bristol N2;
RT "Genome sequence of the nematode C. elegans: a platfor "Genome sequence of the nematode C. elegans: a platfor "Genome sequence of the nematode C. elegans: a platfor "Genome Sequence of the nematode C. elegans: a platfor "Genome Sequence of the nematode C. elegans: a platfor "Genome Sequence of the nematode C. elegans: a platfor "Genome Sequence of the nematode C. elegans: a platfor "Genome Sequence of the nematode C. elegans: a platfor "Genome Sequence of the nematode C. elegans: a platfor "Genome Sequence of the nematode C. elegans: a platfor "Genome Sequence of the nematode C. elegans: a platfor "Genome Sequence of the nematode C. elegans: a platfor "Genome Sequence" of the nematode C. elegans: a platfor "Genome Sequence" of the nematode C. elegans: a platfor "Genome Sequence" of the nematode C. elegans: a platfor "Genome Sequence" of the nematode C. elegans: a platfor "Genome Sequence" of the nematode C. elegans: a platfor "Genome Sequence" of the nematode C. elegans: a platfor "Genome Sequence" of the nematode C. elegans: a platfor "Genome Sequence" of the nematode C. elegans: a platfor "Genome Sequence" of the nematode C. elegans: a platfor "Genome Sequence" of the nematode C. elegans: a platfor "Genome Sequence" of the nematode C. elegans: a platfor "Genome Sequence" of the nematode C. elegans: a platfor "Genome Sequence" of the nematode C. elegans: a platfor "Genome Sequence" of the nematode
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"Molecular characterisation of the Ancylostoma-secreted protein fame from the adult stage of Ancylostoma caninum.";
Int. J. Parasitol. 33:897-907(2003).

EMBL; AY217007; AA063578.1; -; mRNA.
GO; GO:0005576; C:extracellular region; IEA.
InterPro; IPR001283; Allrgn_V5/Tpx1.
Pfam; PF00188; SCP; 2.
SMART; SM00198; SCP; 2.
SMART; SM00198; SCP; 2.
SEQUENCE 451 AA; 48760 MW; 9784247C59C294FB CRC64;
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karyota; Metazoa; Nematoda; Chromadorea; Rhabditida;
cylostomatoidea; Ancylostomatidae; Ancylostomatinae;
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cylostoma caninum
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-OCT-2003 (TrEMBLrel. 25,
creted protein 6.
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114

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Rhabditida;

Rhabditoidea

platform

DB 2;

Length

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PRINTS; PR00838; VSALLERGEN.
SMART; SM00198; SCP; 1.
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SEQUENCE 221 AA; 25494 MW;
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53; Conservative
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Similarity 29.3%;
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  EVIWQEGKACVCDARPDSFCCDNLCDTR
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Search completed: March 3, 2006, 19:09:28
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November 2005

Published_Applications Nucleic Acid and Published_Applications Amino Acid database searches now generate two sets of results each. The Published_Applications databases have been split into two parts to reduce the amount of time required for their daily updates. This results in more machine time being available for processing searches.

Newly published applications will appear in the Published_Applications_New databases; older published applications make up the Published_Applications_Main databases.

Searches run against Nucleic Acid Published_Applications produce two sets of results, with the extensions .rnpbm (Published_Applications_NA_New).

Searches run against Amino Acid Published_Applications produce two sets of results, with the extensions .rapbm (Published_Applications_AA_New).

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Protein Sequence Searches - February 2005

All of the sequence databases on ABSS have recently been updated.

- Please note that the curators of the UniProt database have purged some temporary accession numbers from the most recent version of UniProt. These sequences have been assigned new permanent accession numbers. The new UniProt record may not contain the previous temporary accession number.
- If you encounter an accession number from an older search run against UniProt (results file extension .rup) that can no longer be found in the database, the permanent record with the new accession number can be found by searching the old accession number in the UniProt Protein Archive database (UniPARC) at:

http://www.pir.uniprot.org/database/archive.shtml

If you have any questions regarding this information or your results, please contact any STIC searcher.

When submitting sequence search results for scanning into IFW, please include a copy of this attachment to assist any future Examiners or members of the public who may encounter UniProt temporary accession numbers.

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